8/24/06

T. Gibbs 09/436060 SID 16

SCORE OVER LENGTH SEARCHES

Attached is a score over length search. This search was developed to overcome limitations in most standard search systems which favor large sequences with high scoring, but lesser overall identity over smaller sequences with higher overall identity. This search is especially useful for relatively small nucleic acid or polypeptide target sequences (antisense, fragments, probes, primers, RNAi, epitopes, haptens, etc.) claimed functionally via a form of hybridization and/or identity language and having defined upper and lower polynucleotide and or polypeptide length limits.

The score over length search is performed by first running the query sequence using examiner-specified identity and polynucleotide or protein length limit parameters, and saving 65,000 hits and 0 alignments from each desired database. The resulting output is reformatted using a Microsoft Word macro and is imported into Excel. The summary table data are then sorted by the ratio of score of each hit sequence divided by its length and the accession numbers for all hits below the examiner's desired score over length parameters are deleted. The remaining accession numbers are used to pull the corresponding sequences from the databases into subdatabases enriched for good hits and the query sequence is re-run against these subdatabases to yield the final results.

The score over length cutoff for this search is 75.70

Examiner Please Note: This cover sheet should be included when submitting results to be scanned.

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STIC-Biotech/ChemLib

161786

From:

Gibbs, Terra

Sent:

Monday, August 08, 2005 12:05 PM

To: Subject: STIC-Biotech/ChemLib sequence search request...

I have another request for a score over length search:

I need a length limited nucleotide sequence search of SEQ ID NO:16 in USSN 09/436,060, where the returns are rank ordered based on the score over length/ratio as we've discussed. I need the lengths limited to hits between 8 and 80 nucleotides, and I'll take as many hits as you can import into excel (64,000?), and alignments for anything above .75 on the above ratio. Hope this is clear, please call me if it's not. I also need the interference databases searched.

Terra Cotta Gibbs, Ph.D. Art Unit 1635 Remsen Building 2D10 Mailbox 2C18 571-272-0758

STAFF USE ONLY

Searcher:
Searcher Phone: 2Date Searcher Picked up:
Date Completed:
Searcher Prep/Rev. Time:
Online Time:

Type of Search

NA#:_____ AA#:_____
Interference:___ SPDI:___
S/L:___ Oligomer:____
Encode/Transl:____ Text:__
Structure#:____ Text:__
Inventor:___ Litigation:___

Vendors and cost where applicable STN:______DIALOG:_____

QUESTEL/ORBIT:_______
LEXIS/NEXIS:______

SEQUENCE SYSTEM:

WWW/Internet:

Other(Specify):

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n 5.1.6 ; Compugen Ltd.	Search time 2 Seconds (without alignments) 3.144 Million cell updates/sec	aggactcggctcacacatgc 4	, parameters: 590		2	results predicted by chance to have a to the score of the result being printed, of the total score distribution.	.ES Description		ACCESSION: BD225879 ACCESSION: BD225880 ACCESSION: AX019628	ACCESSION: AX019629 ACCESSION: BD225882 ACCESSION: AX019631	ACCESSION:BD225881 ACCESSION:AX019630 ACCESSION:BD225883	ACCESSION: AX019632 ACCESSION: BD225837 ACCESSION: AX019586	ACCESSION: BD225838 ACCESSION: BD225839	ACCESSION: AXOLDS8 ACCESSION: AXOLD588 ACCESSION: AR279616	ACCESSION; AR305067 ACCESSION; BD071082	ACCESSION: AR016054 ACCESSION: AR059215 ACCESSION: AR063829	ACCESSION: AR063832 ACCESSION: AR075526	SSIO1	SSIO) SSIO)	ACCESSION: AR279620 ACCESSION: AR305070 ACCESSION: AR305071
GenCore v Copyright (c) 1993 -	OM nucleic - nucleic search, using sw model Run on: August 24, 2005, 14:20:26;	Title: US-09-436-060A-16 Perfect score: 451 Sequence: 1 999ttgcggaggggggct Scoring table: IDENTITY NUC Gapop 10.0 , Gapext 0.5	Searched: 295 segs, 6972 residues Total number of hits satisfying chosen para	length: 0 Length: 200000000 Minimum Match 0% Maximum Match 100%	Database : rge.subdb:*	Pred. No. is the number of results prescre greater than or equal to the sor and is derived by analysis of the tot.	SUMMARIES * Regult Query No. Score Match Length DB ID	54 12.0 62 1	* 11 11 11	46.2 10.2 51 1 43 9.5 51 1 43 9.5 51 1	41.4 9.2 51 1 41.4 9.2 51 1 38.2 8.5 51 1	38.2 8.5 51 1 36 8.0 38 1	31.2 6.9 38 1	1, 31.2 6.9 38 1 18 31.2 6.9 38 1 19 31 6.9 31 1	20 31 6.9 31 1	22 30 6.7 30 1 23 30 6.7 30 1 24 30 6.7 30 1	25 30 6.7 30 1 26 30 6.7 30 1	27 30 6.7 30 1 28 30 6.7 30 1	29 30 6.7 30 1 30 30 6.7 30 1	30 6.7 30 1 30 6.7 30 1 30 6.7 30 1

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AUTHORS Roninson, I.B. and Grossman, A. TITLE Compositions, methods and kits for identifying naturally occurring TITLE RNA sequences having affinity for RNA-binding proteins JOURNAL Patent: US 6238867-A 16 29-MAY-2001; FEATURES Location/Qualifiers source /organism="unknown" /mol_type="unassigned DNA"	Query Match 12.0%; Score 54; DB 1; Length 62; Best Local Similarity 100.0%; Pred. No. 1.4; Matches 54; Conservative 0; Mismatches 0; Indels 0; Gaps 0;	Oy 1 GGGTTGCGGAGGGTGGGCCTGGGAGGGTGGCCATTTTTGTCTAACCCTA 54	RESULT 2 AR154150 LOCUS CCUS AR154150 66 bp DNA linear PAT 08-AUG-2001 DEFINITION Sequence 15 from patent US 6238867.	Σ	Unclassified. REFERENCE 1 (bases 1 to 66) AUTHORS Rominson, I.B. and Grossman, A. TITLE Compositions, methods and Alts for identifying naturally occurring TITLE RNA semisonces having affiliative for RNA-binding protesing	JOURNAL Patent: US 623867-A 15.29-MAY-2001; FEATURES Location/Qualifiers 166 /organism="unknown" /mol type="unassigned DNA"	Query Match Best Local Similarity 100.0%; Pred. No. 1.5; Matches 54; Conservative 0; Mismatches 0; Indels 0; Gaps 0;	Oy 1 GGGTTGCGGAGGGTGGGAGGGGTGGTGGCCATTTTTTGTCTAACCCTA 54 	RESULT 3 BD225879 LOCUS BD225879 LOCUS BD225879 S1 bp DNA linear PAT 17-JUL-2003 DEFINITION Promoter region of mouse and human telomerase RNA component genes.		(bases 1 to 51) th, W.N. to concer region of mouse and mouter region of mouse and ent: JP 2002509599-A 82 (ICER RESEARCH CAMPAIGN TEC	COMMENT OS Artificial Sequence PN 17 2002509699-A/82 PD 02-APR-2002 PD 02-APR-2002 PP 29-JAN-1999 GB 9801902.9 PR 29-JAN-1999 GB 9801902.9 PI WILLIAM NICOL KEITH PC C12N15/09,A61K31/7105,A61K31/711,A61K35/76,A61K38/00,A61K45/00, PC A61K48/00, PC A61P5/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02 PC ,C12Q1/68//C12N9/12,
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Gaps

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Gaps

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DEFINITION

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RESULT 5 AX019628

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ACCESSION VERSION KEYWORDS

Best Loc Matches

FEATURES

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bp DNA linear PAT 17-JUL-200 human telomerase RNA component genes.
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                                                                                               Promoter regions of the mouse and human telomerase rna component
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH
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Best Local Similarity 94.1%; Pred. No. 4.2;
Matches 48; Conservative 0; Mismatches 3;
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/organism="synthetic construct"
/wol_type="unassigned DNA"
/db_xref="taxon:32630"
/note="Mutant construct"

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Best Local Similarity 94.1%; Pred. No. 4.2;
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TITLE
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C12N15/09,A61K31/7105,A61K31/711,A61K35/76,A61K38/00,A61K45/00, PC
                                                                                                                                                                                                                                                                                                                                                                                                               bD225880 51 bp DNA linear PAT 17-JUL-200:
Promoter region of mouse and human telomerase RNA component genes.
BD225880
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CANCER RESPARCH CAMPAIGN TECHNOLOGY LTD
OS Artificial Sequence
PN JP 2002505699-A/83
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PC A61P35/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02 PC

C1201/68//C12N9/12,

PC (A61K25/76,A61K31:S22),C12N15/00,A61K37/02,C12N5/00 CC

Description of Artificial Sequence: Mutant construct FH Key
                                                                                                                                                                                                                                                       Gaps
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PC (A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00
Description of Artificial Sequence: Mutant construct FH
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                  TTGTCTAACC 51
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                                                                     /organism='Artificial Sequence'
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                                                                                                                                                                                                               Score 46.2; DB 1;
Pred. No. 4.2;
0; Mismatches 3;
                                                                                                                                   /organism="synthetic construct"
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1 (bases 1 to 51)
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Sequence 82 from Patent W09938964.
AX019628 1GI:10043542
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29-JAN-1999 JP 2000529424
29-JAN-1998 GB 9801902,9
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                                                                                                                                                                                                               th 10.2%; Similarity 94.1%; 48; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               synthetic construct
synthetic construct
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                                                          source
                                                                                                                                                                                                            Query Match
Best Local Similarity
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DEFINITION

RESULT 4 BD225880

g

Matches

FEATURES

ACCESSION VERSION KEYWORDS

ORGANISM

SOURCE

AUTHORS TITLE JOURNAL

REFERENCE

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PAT 07-SEP-2000
                                                                         Artificial Sequence

Artificial Sequence

PN JP 2002509699-A/84

PN 02-APM-1099 JP 2000529424

PR 29-JAN-1998 GB 9801902.9

PI WILLIAM NICOL KEITH

PC 212N15/09,A61K31/7105,A61K31/711,A61K35/76,A61K38/00,A61K45/00, PC
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Promoter region of mouse and human telomerase RNA component genes.
BD225883
                         Promoter region of mouse and human telomerase RNA component genes
Patent: JP 2002509699-A 84 02-APR-2002;
CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
                                                                                                                                                                                                                                             PC A61905/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02 PC (21201/68/C12N9/12, PC (21401/68/C12N9/12, PC (21401/68/C12N9/12, PC (21401/68/C12N5/00,A61K31:522),C12N15/00,A61K31/02,C12N5/00 CC Description of Artificial Sequence: Mutant construct FH Key Location/Qualifiers
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GGGTTGCGGAAAAATGGGCCTGGGTAAGGTGGTGGCCATTTTTTGTCTAACC
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Pred. No. 9.4;
0; Mismatches 6; Indele
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llarity 88.2%; Pred. No. 9.4;
Conservative 0; Mismatches 6;
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/note="Mutant construct"
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    .51
    /organism="synthetic co
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ses 45; Conserv
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Best Local
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BD225883
LOCUS
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ACCESSION
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     AUTHORS
TITLE
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AX019630
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TITLE
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KEYWORDS
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PN JP 2002509699-A/85
PD 02-APR-2002
PF 29-JAN-1999 GB 9801902.9
PR 29-JAN-1999 GB 9801902.9
PR 19-JAN-1999 GB 9801902.9
PI WILLIAM NICOL KEITH
PC C12N15/09, A61K31/7105, A61K31/711, A61K35/76, A61K38/00, A61K45/00, PC A61R48/00, A61K45/00, PC A61R98/00, A61K91/21, C12N5/10, C12P21/02 PC
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Promoter region of mouse and human telomerase RNA component genes.
BD225881
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                                                                                                                                                                   C1201/68//C12N9/12,
PC (A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00 CC
Description of Artificial Sequence: Mutant construct FH Key
Location/Qualifiers
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Location/Qualifiers
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                  9.5%; Score 43; DB 1; Length 51;
larity 90.2%; Pred. No. 7.2;
Conservative 0; Mismatches 5; Indels
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synthetic construct
synthetic construct
other sequences, artificial sequences.
1 (bases 1 to 51)
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Sequence 85 from Patent WO9938964.
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synthetic construct
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Matches 46; Conserv
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BD225881
LOCUS
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VERSION
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SOURCE
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AUTHORS
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DNA linear PAT 17-JUL-200 telomerase RNA component genes.

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Promoter region of mouse and human telomerase RNA component genes Patent: JP 200259659-A 40 02-APR-2002; CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD CO-APR-2002 PD 02-APR-1099 JP 2000529424 PR 29-JAN-1998 GB 9801902.9 PI WILLIAM NICOL KEITH PR 29-JAN-1998 GB 9801902.9 PC CI2NIS/09,A61K31/7105,A61K31/711,A61K35/76,A61K38/00,A61K45/00, PC
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PC A61P55/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02
PC A61P55/00,C12N9/12,
PC (A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00 CC
Description of Artificial Sequence:Oligonucleotide FH Key
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism='Artificial Sequence'
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1. 38
/organism="synthetic construct"
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                                                                                                                                  other sequences; artificial sequences.

(bases 1 to 38)
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         Promoter region of mouse and human BD225837
                                                               BD225837.1 GI:33035607
JP 2002509699-A/40.
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synthetic construct
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                                                                                                                                                                     Keith, W.N.
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                                                                                                 Promoter region of mouse and human telomerase RNA component genes
Patent: JP 2002509699-A 86 02-APR-2002;
CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
CA Artificial Sequence
PN JP 2002509699-A/86
PD 02-APR-2002
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PC A61P35/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02 PC
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PC (A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00
Description of Artificial Sequence: Mutant construct FH
Location/Qualifiers
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/organism='Artificial Sequence'.
Location/Qualifiers
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH
Location/Qualifiers
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/organism="synthetic construct"
/mol_type="genomic DNA"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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                                                other sequences, artificial sequences.
1 (bases 1 to 51)
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Pred. No. 16;
0; Mismatches
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larity 84.3%; Pred. No. 16;
Conservative 0; Mismatches
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Sequence 86 from Patent W09938964.
AX019632
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29-JAN-1998 GB 9801902.9
WILLIAM NICOL KEITH
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illarity 84.3%;
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DB 1; Length 38; 17;

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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH
Location/Qualifiers
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  linear
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                                                                                                                                                                                                                                                                                                                                                                      /organism="synthetic construct"
/mol_type="unassigned DNA"
/mole="taxon:32630"
/noFe="Oligonucleotide"
  DNA
                                                                                                                 synthetic construct
synthetic construct
other sequences; artificial sequences.
AX019586 38 bp
Sequence 40 from Patent WO9938964.
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PAT 07-SEP-2000
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                                       PC A6195/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02 PC (21201/68//C12N9/12, PC (A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00 CC Description of Artificial Sequence:Oligonucleotide FH Key Location/Qualifiers
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
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Patent: WO 9938964-A 42 05-AUG-1999;
KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Localion/Qualifiers
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/organism="synthetic construct"
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/db_kref="taxon:32630"
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llarity 91.7%; Pred. No. 36;
Conservative 0; Mismatches
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6.9%; Score 31.2; D
Best Local Similarity 91.7%; Pred. No. 36;
Matches 33; Conservative 0; Mismatches
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CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
OS Artificial Sequence
PN JP 2005509699-A/41
PD 02-APR-2002
PP 29-JAN-1999 JP 2000529424
PR 29-JAN-1999 GB 9801902.9
PR 29-JAN-1999 GB PRITH
PR 29-JAN-1999 GB PR 29-JAN-1999 GB PRITH
PR 29-JAN-1999 GB PR 29-JAN-199
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                                                                                                                                                       ยบระวย38
Promoter region of mouse and human telomerase RNA component genes.
BD225838
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CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
S Artificial Sequence
NO 3D 2002509699-A/42
PD 02-APR-2002
PF 29-JAN-1999 JP 2000529424
PF 29-JAN-1998 GB 9801902.9
PI WILLIAM NICOL KEITH
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PC A61P35/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02 PC (A61P35/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02 PC (A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00 CC Description of Artificial Sequence:Oligonucleotide FH Key Location/Qualifiers
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Location/Qualifiers
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             GGGTTGCGGAGGGTGGCCTGGAAGGGGTGGTGGCC 38
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/organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
                                                                                                                                                                                                                                                                                                                                         other sequences, artificial sequences.
1 (bases 1 to 38)
Keith, W.N.
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other sequences; artificial sequences.
1 (bases 1 to 38)
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1 Similarity 91.7%; Pred. No. 36;
33; Conservative 0; Mismatches
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bu071082 31 bp RNA linear PAT 27-AUG-2002
Modulation of mammalian telomerase by peptide nucleic acids.
BD071082
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Location/Qualifiers
                                                                                                                    1 (bases 1 to 31)
Shay, J.W., Wright, W.E., Piatyszek, M.A., Corey, D. and Norton, J.C.
Modulation of mammalian telomerase by peptide nucleic acids
Patent: JP 2001517929-A 48 09-OCT-2001;
GERON CORP
                                                                                                                                                                                                   Unidentified
JP 2001517929-A/48
09-OCT-2001
09-APR-1997 JP 1997536487
09-APR-1996 US 08/630019
JERRY W SHAY, WOODRING E WRIGHT, MIECZYSLAW A PIATYSZEK, DAVID
                                                                                                                                                                                                                                                                                                                                                                                                     /organism='Unidentified'
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Strandedness: Single;
Topology: Linear;
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                                                        BD071082.1 GI:22616685 JP 2001517929-A/48. unidentified
                                                                                                                                                                                                                                                                                           COREY,
JAMES C NORTON
                                                                                                 unidentified
unclassified.
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Weinrich, S.L., Atkinson, E.M. III, Lichtsteiner, S.P., Vasserot, A.P.
and Pruzan, R.A.
Purified telomerase
Patent: US 6517834-A 1 11-FEB-2003;
Location/Qualifiers
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Unclassified.
1 (bases 1 to 31)
Weinrich,S.L., Atkinson,E.M. III, Lichtsteiner,S.P., Vasserot,A.P. and Pruzan,R.A.
Methods for purifying telomerase
Patent: US 6545133-A.
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                                                                          DB 1;
 /organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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                                                                       Score 31.2; DB; Pred. No. 36; 0; Mismatches
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Sequence 1 from patent US 6545133.
AR305067
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/organism="unknown"
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                                                                         ch 6.9%;
1 Similarity 91.7%;
33; Conservative
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AR279616/c
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AR305067/c
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Unclassified.
1 (bases 1 to 30)
1 (laponteau, B., Feng, J., Funk, W. and Andrews, W.H. Assays for the DNA component of human telomerase Patent: US 5776679-A 22 07-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                Length 30;
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1 TTTTGTCTAACCCTAACTGAGAAGGGCGTAG 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        77 IGCTITIGGICCCCGCGCGCTGTTTTCTC 106
                                                                                                      AR016054 30 bp
Sequence 22 from patent US 5776679.
AR016054 GI:3972331
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1 (bases 1 to 30)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomorase
Patent: US 6258535-A 22 10-JUL-2001;
                                                                                                                                                                                                                                                                     1 (bases 1 to 30)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5958680-A 23 28-SEP-1999;
          Length 30;
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                                                               137 CCTGCCGCCTTCCACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                         77 TGCTTTTGCTCCCGCGCGCTGTTTTCTC 106
        Query Match 6.7%; Score 30; DB Best Local Similarity 100.0%; Pred. No. 34; Matches 30; Conservative 0; Mismatches
                                                                                                                                                       30 bp
Sequence 23 from patent US 5958680.
AR075526.1 GI:10002274
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                                                                             CCTGCCGCCTTCCACCGTTCATTCTAGAGC 1
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Mammalian telomerase.
BD176165
BD176165.1 GI:29121871
JP 2002272489-A/24.
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Unclassified.
1 (bases 1 to 30)
Kim,N.Woo., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.
Methods for detecting the RNA component of telomerase
Patent: US 5846723-A 8 08-DEC-1998;
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Unknown.
Unclassified.
Unclassified.
1 (bases 1 to 30)
Kim,N.Woo., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.
Methods for detecting the RNA component of telomerase
Patent: US 5846723-A 5 08-DEC-1998;
Location/Qualifiers
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(1 (bases 1 to 30)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5837857-A 22 17-NOV-1998;
Locarion/Qualifiers
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           AR059215
Sequence 22 from patent US 5837857.
AR059215.1 GI:5984792
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Sequence 5 from patent US 5846723.
AR063829.1 GI:5993137
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Sequence 8 from patent US 5846723.
AR063832.1 GI:5993140
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Best Local Similarity 100.09
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AR063832/c
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AR063829/c
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PAT 10-APR-2003

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Unknown.
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Unclassified.
Weinsich,S.L., Atkinson,E.M. III, Lichtsteiner,S.P., Vasserot,A.P. and Pruzan,R.A.
Purified telomerase
Purified telomerase
Patent: US 651734.A 4 11-FEB-2003;
Location/Qualifiers
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1 (bases 1 to 30)
Weinrich,S.L., Atkinson,E.M. III, Lichtsteiner,S.P., Vasserot,A.P. and Pruzan,R.A.
Purified telomerase
Patent: US 6517834-A 5 11-FEB-2003;
Location/Qualifiers
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Weinrich, S.L., Atkinson, E.M. III, Lichtsteiner, S.P., Vasserot, A.P. and Pruzan, R.A.
Methods for purifying telomerase
Patent: US 6545133-A 4 08-APR-2003;
Location/Qualifiers
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Sequence 5 from patent US 6517834.
AR279620 GI:29714514
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Sequence 4 from patent US 6545133.
AR305070.
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AR305070/c
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PN 24 2002272489-A/24

PP 05 4-8ZEP-2002

PR 06-MAR-2002

PR 07-JUL-1994 US 08/272102,27-OCT-1994 US 08/330123 PR

O7-JUL-1995 US 08/472802,07-JUN-1995 US 08/482115 PI BRYANT

VILLEDONTEAL JUNIA FENG, WALTER FINK, WILLIAM HANDREWS PC

C12N15/09,C12N9/99,C12Q1/68,G01N33/566,C12N15/00 CC

Strandedness: Single,

CC Topology: Linear;

CC Mammalian telomerase

FH Key Location/Qualifiers

FT source location/Qualifiers

FT Source /Organism='Unidentified'.
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Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: 7P 2002272489-A 24 24-SEP-2002;
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Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: US 5583016-A 22 10-DEC-1996;
Location/Qualifiers
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AR279619
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RESULT 29

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LOCUS DEFINITION ACCESSION VERSION KEYWORDS SOURCE

RESULT 30 AR279619/c

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PAT 16-JUL-2002

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BD023701 30 bp DNA linear PAT 27-AUG-2002 Method for detecting and inhibiting RNA component of telomerase.
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1 (bases 1 to 30)

1 (kim,N. Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L. Method for detecting and inhibiting RNA component of telomerase Patent: JP 2001507229-A 5 05-JUN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PN JP 2001507229-A/5
PD 05-JUN-2001
PF 19-DEC-1997 JP 1998529003
PR 20-DEC-1997 US 08/770564, 20-DEC-1996 US 08/770565
NAM WOO KIM, FRED WU, JAMES T KEALEY, RONALD PRUZAN, SCOTT L PI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Letsinger,R.L. and Garimella,V.
Method of detection by enhancement of silver staining
Patent: WO 0204681-A 2 17-JAN-2002;
NORTHWENERN UNIVERSITY (US)
Location/Qualifiers
1 (bases 1 to 30)
Letsinger.R.L. and Garimella,V.
Method of detection by enhancement of silver staining
Patent: 10.5 6602669-A 2 05-AUG-2003;
Location/Qualifiers
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Sequence 2 from Patent WO0204681.
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BD023701.1 GI:22564924
JP 2001507229-A/5.
unidentified
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AX465471/c
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Weinrich, S.L., Atkinson, E.M. III, Lichtsteiner, S.P., Vasserot, A.P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: US 6548296-A 22 15-APR-2003;
Location/Qualifiers
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34;
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                   137 CCTGCCGCCTTCCACCGTTCATTCTAGAGC 166
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     6.7%; Score 30; DB 100.0%; Pred. No. 34; ive 0; Mismatches
                                                                                                                                                                                                                                                                                                               and Pruzan, R.A.
Methods for purifying telomerase
Patent: US 654513-A 5 09-APR-2003;
Location/Qualifiers
                                                                                                                                    30 bp
Sequence 5 from patent US 6545133.
AR305071.1 GI:31694378
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Sequence 22 from patent US 6548298.
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Sequence 2 from patent US 6602669.
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    /organism="unknown"
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AR306472.1 GI:31696311
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Best Local Similarity 100.
Matches 30; Conservative
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Unclassified.
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Unclassified.
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RESULT 34 AR306472/c

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DEFINITION

ACCESSION VERSION KEYWORDS

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REFERENCE AUTHORS source

TITLE JOURNAL FEATURES

RESULT 35 AR373060/c

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PAT 21-JAN-2000

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Unclassified.
Unclassified.
1 (bases 1 to 30)
Weinirch,S.L., Atkinson,E.M. III, Lichtsteiner,S.P., Vasserot,A.P.,
Pruzan,R.A. and Realey,J.T.
Purified telomerase
Patent: US 5968506-A 2 19-OCT-1999,
Location/Qualifiers
1 . .30
/organism="unknown"
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Best Local Similarity 100.0%; Pred. No. 40;
Matches 29; Conservative 0; Mismatches (
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 Atkinson, E.M. and Kealey, J.T.
PURIFIED TELOMERASE
Patent: WO 9845450-A 5 15-OCT-1998;
GERON CORP (US)
                                                                                                          /organism="unidentified"
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/db_xref="taxon:32644"
                                                                                                                                                                                                                                                                                                            167 AAACAAAAATGTCAGCTGCTGGCCCGTT 195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Patent: WO 9845450-A 6 15-OCT-1998;
GERON CORP (US)
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/mod_base=OTHER
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Squence 2 from patent US 5968506.
AR079889.1 GI:10006642
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Sequence 6 from Patent WO9845450.
A84596.1 GI:6733512
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/organism="unidentified"
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Atkinson, E.M. and Kealey, J.T.
PURIFIED TELOMERASE
                                                                            Location/Qualifiers
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AR079889/c
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A84596/c
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PN JP 201507229-A/8
PD 05-JUN-2001
PF 19-DEC-1997 JP 1998529003
PR 20-DEC-1996 US 08/770564, 20-DEC-1996 US 08/770565 PI
NAM WOO KIM, FRED WU, JAMES T KEALEY, RONALD PRUZAN, SCOTT L PI
WEINRICH
PC C12N19/10, AG1K9/08, AG1K31/7105, AG1K45/00, AG1F48/00, AG1P35/00, PC C12N19/10, DC C12N19/10, PC POPOLOGY: Linear;
CC /note= '0ligo 21'
FH KEY
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Method for detecting and inhibiting RNA component of telomerase.
BD023704
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Kim_N.W., Wu,F., Kealey,J.T., Fruzan,R. and Weinrich,S.L.
Method for detecting and inhibiting RNA component of telomerase
Patent: JP 2001507229-A 8 05-JUN-2001;
GERON CORP
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6.7%; Score 30; DB
Best Local Similarity 100.0%; Pred. No. 34;
Matches 30; Conservative 0; Mismatches
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Sequence 5 from Patent WO9845450.
A84595
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JP 2001507229-A/8.
unidentified
unidentified
Strandedness: Single;
CC Topology: Linear;
CC /note= 'oligo 16'
FH Key
                      Topology: Linear;
/note= 'oligo 16'
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Best Local Similarity 100.
Matches 30; Conservative
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1 (bases 1 to 30)
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BD023704/c
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AUTHORS TITLE

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Gaps

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PAT 31-AUG-2000

ORGANISM

SOURCE

REFERENCE

ACCESSION VERSION KEYWORDS

RESULT 39 A84595/c

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US09436060A-16.rge.sl

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Examays

Clade; Panicoideae; Andropogoneae; Eas.

Clade; Panicoideae; Andropogoneae; Zea.

Clade; Panicoideae; Andropogoneae; Zea.

Clade; Panicoideae; Andropogoneae; Zea.

Examan, R.A. and Kealey, J.T.

Prizan, R.A. and Kealey, J.T.

Prizan, R.A. and Kealey, J.T.

Purified telomerase

AL Patent: JP 2001509681-A 6 24-JUL-2001;

GERON CORP

PN JP 2001509681-A/6

PD 24-JUL-2001;

GERON CORP

PR 04-APR-1997 JP 1998542718

PI SCOTT L WEINRICH; EDWARD M ATKINSON III, SERGE P LICHTSTEINER,

PI ALAIN P VASSEROT, ROMALD A PRUZAN, JAMES T KEALEY PC

CI2N15/54, CI2N9/12, CO7K16/40, C12Q1/68, C07K14/47 CC Strandednese:
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                                                                        Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Viridiplantae; Streptophyta; Endracophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD clade; Panicoideae; Andropogoneae; Zea.
1 (bases 1 to 30)
Weinrich, S.L., Iii, E.M.A., Lichtsteiner, S.P., Vasserot, A.P., Pruzan, R.A., and Kealey, J.T.
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/mod_base= OTHER
/note= 'N = biotinylated G'
/note= 'oligonucleotide 14'
/ov
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6.4%; Score 29; DB
Best Local Similarity 100.0%; Pred. No. 40;
Matches 29; Conservative 0; Mismatches
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              BD058136.1 GI:22603742
JP 2001509681-A/5.
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JP 2001509681-A/6.
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Purified telomerase.
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BD058137/c
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Weinrich, S.L., Atkinson, E.M. III, Lichtsteiner, S.P., Vasserot, A.P., Pruzan, R.A. and Kealey, J.T.
Pruzin, E.A. and Kealey, J.T.
Purified telomeras S968506-A 5 19-OCT-1999;
Location/Qualifiers
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Weinrich, S.L., Arkinson, E.M. III, Lichtsteiner, S.P., Vasserot, A.P., Pruzan, R.A. and Kealey, J.T.
Pruzan, R.A. and Kealey, J.T.
Purified telomerase
Purified telomerase
Patent: Location/Qualifiers
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                              Query Match 6.4%; Score 29; DB 1; Length 30; Best Local Similarity 100.0%; Pred. No. 40; Matches 29; Conservative 0; Mismatches 0; Indels
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/organism="unknown"
/mol_type="unassigned DNA"
 /mol_type="unassigned DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    167 AAACAAAAATGTCAGCTGCTGGCCCGTT 195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           137 CCTGCCGCCTTCCACCGTTCATTCTAGAG 165
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                                                                                             43 TGTCTAACCCTAACTGAGAAGGGCGTAGG 71
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/organism="unknown"
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Purified telomerase.
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Best Local Similarity 100.0
Matches 29; Conservative
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BD058136/c
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AR079893/c
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AR079892/c
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1 (bases 1 to 30)

Weinrich,S.D., Atkinson,E.M. III, Lichtsteiner,S.P., Vasserot,A.P.
Meinrich,S.D., Atkinson,E.M. III, Lichtsteiner,S.P., Vasserot,A.P.
and Pruzan,R.A.
Methods for purifying telomerase
Patent: US 6545133-A 3 08-APR-2003;
Location/Qualifiers

1. .30
       PC
Strandedness:
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1 (bases 1 to 30)
Weinrich, S.L., Atkinson, E.M. III, Lichtsteiner, S.P., Vasserot, A.P.
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PI ALAIN P VASSEROT, RONALD A PRUZAN, JAMES T KEALEY PC C12N15/54, C12N9/12, C07K16/40, C12Q1/68, C07K14/47 CC Strissingle;
CC /mod_base= OTHER
CC /mode= 'N = biotinylated A'
FC /mode= 'N = biotinylated A'
FF KEY
FF Modified base 1
FT modified base 31.
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                                                                                                                                                                                                                                                                                                                DB 1; Length 31;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        and Pruzan, R.A.
Purified telomerase
Patent: US 6517834-A 3 11-FEB-2003;
Location/Qualifiers
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Sequence 3 from patent US 6517834.
AR279618 GI:29714512
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Sequence 3 from patent US 6545133.
AR305069.1 GI:31694376
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/db_xref="taxon:4577"
                                                                                                                                                                                                                                                 /organism="Zea mays"
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AR305069/c
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AR279618/c
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Spermatophyta, Magnoliophyta, Liliopsida, Poales, Poaceae, PACCAD
clade, Panicoideae, Andropogoneae, Zea.
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24-JUL-2011
04-APR-1997 JP 1998542718
SCOTT L WEINRICH, EDWARD M ATKINSON III, SERGE P LICHTSTEINER,
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Weinrich, S.L., Iii, E.M.A., Lichtsteiner, S.P., Vasserot, A.P., Pruzan, R.A. and Kealey, J.T.
Purified telomerase
Patent: JP 2001509681-A 2 24-JUL-2001;
GERON CORP
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100.0%; Pred. No. 42;
tive 0; Mismatches 0; Indels
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                                                                                                 6.4%; Scc...
100.0%; Pred. No. .c.,
0, Mismatches
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GERON CORP (US)

    .31
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    /mol_type="unassigned DNA"
    /db_xref="taxon:32644"

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/mod_base=OTHER
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                                                                                                                                                                                                                                                                                               31 bp
Sequence 2 from Patent WO9845450.
A84592
A84592.1 GI:6733508
                          1. .30
/organism="Zea mays"
/mol_type="genomic DNA"
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Atkinson, E.M. and Kealey, J.T.
PURIFIED TELOMERASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
          Location/Qualifiers
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BD058133.1 GI:22603739
JP 2001509681-A/2.
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A84592/c
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GERON CORP

PN 472 201509581-A/4

PD 24-JUL-2001 PF 998542718

PI SCOTT L WEINRICH, EDWARD M ATKINSON III, SERGE P LICHTSTEINER, PI ALAIN P VASSEROT, RONALD A PRUZAN, JAMES T KEALEY PC (12N15/54,C12N9/12,C07K16/40,C12Q1/68,C07K14/47 CC Strandedness: Single; CC /mod base= OTHER CC /mod base= OTHER CC /note= 'oligonucleotide 5'

FH Key Location/Qualifiers

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                               PAT 31-AUG-2000
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Weinrich,S.L., Atkinson,E.M. III, Lichtsteiner,S.P., Vasserot,A.P.,
Pruzan,R.A. and Kealey,J.T.
Purified telomerase
Parent: US 5968506-A 4 19-OCT-1999;
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta; Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD clade; Panicoideae; Andropogoneae; Zea.

1 (bases 1 to 30)
Weinrich, S.L., Iii, E.M.A., Lichtsteiner, S.P., Vasserot, A.P., Purzan, R.A. and Kealey, J.T.
Purified telomerase
Patent: JP 2001509681-A 4 24-JUL-2001;
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                                                                                                                                                                                                                                                                                                        6.1%; Score 27.4; D
96.6%; Pred. No. 52;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                     412 GAGCTGTGGGACGTGCACCCAGGACTCGG 440
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                         30 bp
Sequence 4 from patent US 5968506.
AR079891
                                                                                                                                                                                                                                                                         /mol_type="unassigned DNA
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/organism="Zea mays"
/mol_type="genomic DNA"
/db_xref="taxon:4577"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30 bp
                                                                                                                                                                                                                                                             /organism="unknown"
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JP 2001509681-A/4.
Zea mays
Zea mays
                                                                          AR079891.1 GI:10006644
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Purified telomerase.
BD058135
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Best Local Similarity 96.6'
Matches 28; Conservative
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Unclassified.
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Matches 28; Conserv
                                                                                                      Unknown.
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AR079891/c
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BD058135/c
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                                               DB 1; Length 30;
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                                             Score 28.4; DE Pred. No. 45; 0; Mismatches
                                                                                                        412 GAGCTGTGGGACGTGCACCCAGGACTCGGC 441
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 6.2%; Score 28; DB Best Local Similarity 100.0%; Pred. No. 44; Matches 28; Conservative 0; Mismatches
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Atkinson, E.M. and Kealey, J.T.
PURIFIED TELOMERASE
Batent: WO 9845450-A 4 15-OCT-1998;
GERON CORP (US)
                                                                                                                                                                                                                                                                                                                                      1 (bases 1 to 28)
Greider, C.W. and Le, S.
Telomerase-associated proteins
Patent: US 6300131-A 3 09-OCT-2001;
Location/Qualifiers
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/db_xref="taxon:32644"
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/mod_base=OTHER
                                                                                                                                 30 GAGCTATGGGACGTGCACCCAGGACTCGGC 1
                                                                                                                                                                                                          Sequence 3 from patent US 6300131.
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/organism="unidentified"
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Sequence 4 from Patent W09845450.
A84594
 /organism="unknown"
/mol_type="genomic DNA"
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/organism="unknown"
/mol_type="genomic DNA"
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                                            ch 6.3%;
1 Similarity 96.7%;
29; Conservative
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A84594/c
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AL Patent: JP 2001507229-A 26 05-JUN-2001;
GERON CORP
PN JP 2001507229-A/26
PD 05-JUN-2001
PF 19-DEC-1997 JP 1998529003
PR 20-DEC-1996 US 08/770565 PI
NAM WOO KIM,FRED WU,JAMES T KEALEY,RONALD PRUZAN,SCOTT L PI
WEINRICH
PC C12NN5/10, A61K9/08,A61K31/7105,A61K45/00,A61P35/00,PC
C12NN5/10,C1201/68,C1201/68,C12N15/00,C12N5/00 CC
Strandedness: Single
CC Topology: Linear;
CC Topology: Line
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CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
OS Artificial Sequence
RN JP 2002509699-A/4
PD 02-APR-2002
PF 29-JAN-1999 JP 2000529424
PR 29-JAN-1999 GB 9801902.9
PI WILLIAM NICOL KEITH
PC C12N15/09,A61K31/7105,A61K31/711,A61K35/76,A61K38/00,A61K45/00, PC
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Promoter region of mouse and human telomerase RNA component genes.
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PC A61P35/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02
C12Q1/68//C12N9/12,
PC (A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00 CC
Description of Artificial Sequence: Primer
FH Key Location/Qualifiers
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/organism="synthetic construct"
/mol_type="genomic DNA"
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JP 2002509699-A/4.
synthetic construct
synthetic construct
other sequences; artificial sequences.
1 (bases 1 to 30)
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BD225801/c
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Kim,N.W., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.
Method for detecting and inhibiting RNA component of telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chen, X.Q. and Anker, P.
Cancer diagnosis method
Patent: WO 0190409-A 2 29-NOV-2001;
Chen, Xu Qi (US) ; Stroun, Maurice (CH) ; Anker, Philippe (CH)
Location/Qualifiers
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1 (bases 1 to 27)

1 (hases 1 to 27)

1 (kaley,J.T., Pruzan,R. and Weinrich,S.L. Methods for detecting the RNA component of telomerase Patent: US 5846723-A 26 08-DEC-1998;
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6.0%; Score 27; DB 1; Length 27;
Best Local Similarity 100.0%; Pred. No. 50;
Matches 27; Conservative 0; Mismatches 0; Indels
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/organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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                                                                                                                                 Sequence 26 from patent US 5846723.
AR063850
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/mol_type="unassigned DNA"
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Vallaponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Wammanlian telomerase
Patent: US 5958680-A 38 28-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      unidentified unclassified.

1 (bases 1 to 28)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent 'PP 2002272489-A 33 24-SEP-2002;
GERON CORP
                              Andrews, W.H.
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1 (bases 1 to 28)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H Assays for the DNA component of human telomerase Patent: US 5776679-A 40 07-JUL-1998;
Location/Qualifiers
                                                                                                                                                                                                                                       Query Match 5.9%; Score 26.4; DB 1; Best Local Similarity 96.4%; Pred. No. 57; Matches 27; Conservative 0; Mismatches 1;
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                                                                                                                                                            /organism="unknown"
/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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JP 2002272489-A/33.
unidentified
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BD176174
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
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Unclassified.

1 (bases 1 to 28)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Assays for the DNA component of human telomerase Patent: US 5776679-A 31 07-JUL-1998;
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|organism="synthetic construct"
|mol_type="unassigned DNA"
|db_xref="teaxon:32630"
|note="primer"
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other sequences; artificial sequences.
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Best Local Similarity 93.3%; Pred. No. 58;
Matches 28; Conservative 0; Mismatches
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5.9%; Score 26.4; D
Best Local Similarity 96.4%; Pred. No. 57;
Matches 27; Conservative 0; Mismatches
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/organism="unknown"
/mol_type="unassigned DNA"
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  30 CTAACCCTAACTGAGAAGGGCGTAGGATCC 1
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Sequence 4 from Patent WO9938964.
AX019550
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Unclassified.
                                                                                                                                                                                                                                                                                                                                                                        Keith, W.N.
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DEFINITION ACCESSION VERSION KEYWORDS

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1 (bases 1 to 26)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Assays for the DNA component of human telomerase Patent: US 5776679-A 23 07-JUL-1998;
Location/Qualifiers
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(bases 1 to 26)

(illeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Assays for the DNA component of human telomerase Patent: US 5776679-A 29 07-JUL-1998;

Location/Qualifiers
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5.8%; Score 26; DB 1; Length 26;
Best Local Similarity 100.0%; Pred. No. 56;
Matches 26; Conservative 0; Mismatches 0; Indels
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/organism="unknown"
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Sequence 25 from patent US 5858777.
AR028785 1 GI:5940758
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Sequence 23 from patent US 5776679.
AR016055
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Sequence 29 from patent US 5776679.
AR016061.1 GI:3972338
                145 CTTCCACCGTTCATTCTAGAGCAAAC 170
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                                  26 CTTCCACCGTTCATTCTAGAGCAAAC 1
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1 (bases 1 to 26)
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AR028785
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Emrich, T.D.
Method for detection of carcinoma of the urinary bladder within a urine sample sample Patent: EP 0926245-A 1 30-JUN-1999;
ROCHE DIAGNOSTICS GMBH (DE)
Location/Qualifiers
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/organism='Unidentified'.
Location/Qualifiers
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    .26
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    /db_xref="taxon:32644"

    .26
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urine sample
Patent: EP 0926245-A 2 30-JUN-1999;
ROCHE DIAGNOSTICS GMBH (DE)
Location/Qualifiers
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    .28
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    /db_xref="taxon:32644"

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Sequence 1 from Patent EP0926245.
A94987
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Best Local Similarity 96.4
Matches 27; Conservative
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A94987
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A94988/c
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                                                                                                                                                             Unclassified.

1 (bases 1 to 26)

Kim,N.Woo., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L. Methods for detecting the RNA component of telomerase Patent: US 5846723-A 25 08-DEC-1998;

Location/Qualifiers
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UnClassified.
1 (bases 1 to 26)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Willeponteal an telomerase
Patent: US 5958680-A 24 28-SEP-1999;
Location/Qualifiers
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Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5958680-A 30 28-SEP-1999;
Location/Qualifiers
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.00.0%; Pred. No. 56;
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Sequence 24 from patent US 5958680.
AR075527
AR075527.1 GI:10002275
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AR075533
AR075533.1 GI:10002281
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Sequence 25 from patent US 5846723.
AR063849
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AR075527/c
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Unclassified.
1 (Bessel to 26)
Villepsneau, B., Feng, J., Andrews, W.H. and Adams, R.R.
Methods and reagents for regulating telomere length and telomerase activity.
Villeponteau, B., Feng, J., Andrews, W.H. and Adams, R.R.
Methods and reagents for regulating telomere length and telomerase
activity
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Villaponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5837857-A 23 17-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 5.8%; Score 26; DB 1; Length 26; Best Local Similarity 100.0%; Pred. No. 56; Matches 26; Conservative 0; Mismatches 0; Indels
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Best Local Similarity 100.0%; Prad. No. 56;
Matches 26; Conservative 0; Mismatches
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100.0%; Pred. No. 56;
tive 0; Mismatches
                                            Patent: US 5858777-A 25 12-JAN-1999;
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Sequence 26 from patent US 5858777.
AR028786
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Sequence 23 from patent US 5837857.
AR059216
AR059216.1 GI:5984793
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AR028786/c
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AR059216/c
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Method for detecting bladder cancer in urine samples.
E36507
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                                                                                                                                                                                                                                                                                                  linear
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/organism='Unidentified'
Location/Qualifiers
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56;
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synthetic construct
other sequences; artificial sequences.
1 (bases 1 to 26)
                                                               Query Match 5.8%; Score 26; DB Best Local Similarity 100.0%; Pred. No. 56; Matches 26; Conservative 0; Mismatches
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/organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
                                                                                                                                                   145 CITCCACCGITCATTCTAGAGCAAAC 170
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unidentified
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Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: JP 2002272489-A 25 24-SEP-2002;
GERON CORP
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/organism='Unidentified'.
Location/Qualifiers
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                                            /mol_type="unassigned DNA"
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Sequence 23 from patent US 6258535.
AR161925
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/organism="unknown"
                      /organism="unknown"
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BD176166/c
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C12Q1/02,
C12Q1/48, C12Q1/68, G01N33/15, G01N33/48, G01N33/50//C07K14/47, PC
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C12N15/09,A61K31/70,A61K38/55,A61K39/395,A61K39/395,A61K48/00,
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PC (C12N9/12,C12R1:91),C12N15/00,A61K37/64,C12N5/00 CC
Strandedness: Single;
CC Topology: Linear;
FH Key Location/Qualifiers
FT source / Organism='Unidantifiers'
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unidentified
unclassified.

1 (Dasse 1 to 26)

1 Thomas.R.S. Jochimu.R., Toru.N., Karen,B.C., Greg,B.M.,
Calvin,B.H. and William,H.A.
Human telomerase catalytic eubunit promoter
Detent: JP 1999253177-A 254 21-SEP-1999;
JERON CORP. UNIVERSITY TECHNOLOGY CORP
OS Unidentified
PR 19-05-1999
PP 15-OCT-1999
PP 01-OCT-1999
PR 01
SM unidentified

unclassified.

E 1 (bases 1 to 26)

E Thomas, R.S., Jochimu, R., Toru, N., Karen, B.C., Greg, B.M.,
Calvin, B.H. and William, H.A.
Human telomerase catelytic subunit promoter
Human telomerase catelytic subunit promoter
Patent: JP 1999253177-A 253 21-SEP-1999;
Os Unidentified
PN JP 1999253177-A/253
PD 21-SEP-1999
PF 15-OCT-1999 JP 1998320169
PF 15-OCT-1996 US 08/46.017, 06-MAY-1997 US 08/844.419,
25-APR-1997 US 08/46.017, 06-MAY-1997 US 08/851.843, PR
14-AUG-1997 US 08/46.017, 06-MAY-1997 US 08/911.312, PR
14-AUG-1997 US 08/912.951, 14-AUG-1997 US 08/911.312, PR
R SECHI, JOCHIMU RINGNER, TORU NARAMURA, KAREN B CHAPMAN, PI GR
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08/911.312, PR
08/915.503 PI
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/organism='Unidentified'
Location/Qualifiers
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Human telomerase catalytic subunit promoter.
E37046

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    /organism="unidentified"

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/db_xref="taxon:32644"
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JP 1999253177-A/254.
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Method for detecting bladder cancer in urine samples.
836508
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OS Artificial Sequence
PN UP 199943995-A/2
                                                                                                                                                                                                                                                                    ce | 0.26 | /organism='Artificial Sequence'
Location/Qualifiers | 1.26 | /organism="synthetic construct" | /mol_type="genomic DNA" | /db_xref="taxon:32630"
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Human telomerase catalytic subunit promoter.
E37045.
E37045.1 GI:13023008
JP 1999253177-A/253.
unidentified
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/organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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synthetic construct
other sequences; artificial sequences.
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100.0%; Pred. No. 56;
tive 0; Mismatches
       14-SEP-1999
22-DEC-1998 JP 1998365689
22-DEC-1997 DE 19757300:2
THOMAS ENRIHI
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22-DEC-1998 JP 1998365689
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THOMAS ENRIHI
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JP 1999243995-A/2.
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1 (bases 1 to 26)
Cech, T.R., Lingner, J., Nakamura, T., Chapman, K.B., Morin, G.B.,
Harley, C.B. and Andrews, W.H.
Human telomerase catalytic subunit: diagnostic and therapeutic
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Patent: US 6548298-A 23 15-APR-2003;
Location/Qualifiers
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Location/Qualifiers
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Sequence 312 from patent US 6475789.
AR245519 1 GI:27290730
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Sequence 23 from patent US 6548298.
AR306473.1 GI:31696312
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/organism="unknown"
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AR243519/c
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C12Q1/48, C12Q1/68, G01N33/15, G01N33/48, G01N33/50//C07K14/47, PC
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CALVIN B HAREI,WILLIAM H ANDREWS
CI2N15/09,A61K31/70,A61K38/55,A61K39/395,A61K39/395,A61K48/00,
                                                                                                                                                PC (C12N9/12, C12R1:19), (C12N9/12, C12R1:19), (C12N9/12, C12R1:84), PC (C12N9/12, C12R1:91), C12N15/00, A61K37/64, C12N5/00 CC Strandedness: Single; CC Topology: Linear; Location/Qualifiers FT Source Location/Qualifiers FT Source / Organism='Unidentified'
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Cech.T.R., Lingmer.J., Nakamura,T., Chapman,K.B., Morin,G.B.,
Harley,C.B. and Andrews,W.H.
Human telomerase catalytic subunit: diagnostic and therapeutic
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Sequence 311 from patent US 6475789.
AR243518.1 GI:27290729
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Sequence 23 from patent US 5583016.
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/mol_type="unassigned DNA"

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    /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"

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Kopreski,M.S. and Gocke,C.D.
Method for detection of hTR and hTERT telomerase-associated RNA in
plasma or serum
Patent: US 6607888-A 4 19-AUG-2003;
Location/Qualifiers
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Best Local Similarity 100.0%; Pred. No. 56;
Matches 26; Conservative 0; Mismatches
                                           Query Match 5.8%; Score 26; DB Best Local Similarity 100.0%; Pred. No. 56; Matches 26; Conservative 0; Mismatches
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Greider, C.W. and Le, S.
Telomerase-associated proteins
Patent: US 6300131-A 4 09-OCT-2001;
Location/Qualifiers
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Sequence 4 from patent US 6300131.
AR370169
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Sequence 5 from patent US 6607898.
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/organism="unknown"
/wol_type="genomic DNA"
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Unknown.
Unclassified.
I (bases 1 to 26)
Villeponteau.B., Feng,J., Andrews,W.H. and Adams,R.R.
Villeponteau.B., Feng,J., Andrews,W.H. and Adams,R.R.
Peptides related to TPC2 and TPC3, two proteins that are coexpressed with telomerase activity
Patent: US 6300110-A 25 09-0CT-2001;
Location/Qualifiers
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I (bases 1 to 26)

Villeponteau B., Feng, J., Andrews, W.H. and Adams, R.R.

Peptides related to TPC2 and TPC3, two proteins that are coexpressed with telomerase activity

Patent: US 6300110-A 26 09-0CT-2001;

Location/Qualifiers
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Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 6548298-A 30 15-APR-2003;
Location/Qualifiers
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Sequence 30 from patent US 6548298.
AR306480
AR306480.1 GI:31696319
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AR369723
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/mol_type="genomic DNA"

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1 (bases 1 to 26)
Cech, T.R., Lingner, J., Nakamura, T., Chapman, K.B., Morin, G.B.,
Harley, C.B. and Andrews, W.H.
Cells immortalized with telomerase reverse transcriptase for use in
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(bases 1 to 26)

(cch,T.R., Lingmer,J., Nakamura,T., Chapman,K.B., Morin,G.B.,
Harley,C.B. and Andrews,W.H.

Cells immortalized with telomerase reverse transcriptase for use
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Patent: US 6617110-A 598 09-SEP-2003;
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Sequence 597 from patent US 6617110.
AR393334
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Sequence 598 from patent US 6617110.
AR393335.1 GI:40118740
Best Local Similarity 100.0%; Pred. No. 56; Matches 0; Mismatches
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Sequence 25 from Patent EP0953042.
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/mol_type="genomic DNA"
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Unclassified.
1 (bases 1 to 26)
1 (bases 1 to 26)
Kopreski,M.S. and Gocke,C.D.
Method for detection of hTR and hTERT telomerase-associated RNA in plasma or serum
Patent: US 6607898-A 5 19-AUG-2003;
Location/Qualifiers
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Seguence 597 from patent US 6610839.
AR390720
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Sequence 598 from patent US 6610839.
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             AR381130.1 GI:40088891
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Cancer diagnosis method
Patent: WO 0190409-A 1 29-NOV-2001;
Chen, Xu Qi (US) ; Stroun, Maurice (CH) ; Anker, Philippe (CH)
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Detecting telomerase activity
Patent: WO 0046601-A 9 10-AUG-2000;
LARSEN FRANK (NO); SKAANSENG MARIANNE (NO)
LOCATION/Qualifiers
 1. .26
/organism="synthetic construct"
/mol type="unassigned DNA"
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/db_xref="taxon:32630"
/note="PCR primer"
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/db_xref="taxon:32630"
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Sequence 1 from Patent WO0190409.
AX317988
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Sequence 9 from Patent WO0046601.
AX033377
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                                                                   /note="PCR primer"
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                                                                                              Andrews, W.H., Villeponteau, B., Adams, R.R. and Feng, J. Methods and reagents for regulating telomere length and telomerase activity Patent: FP 0953042-A 25 03-NOV-1999; GERON CORP (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Andrews, W.H., Villeponteau, B., Adams, R.R. and Feng, J.
Methods and reagents for regulating telomere length and telomerase
activity
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Detecting telomerase activity
Patent: WO 0046601-A 8 10-AUG-2000;
LARSEN FRANK (NO); SYGANSENG MARIANNE (NO)
LOCATION/Qualifiers
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    .26
    /organism="unidentified"
    /mol_type="unassigned DNA"
    /db_xref="taxon:32644"

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GERON CORP (US)

    .26
/organism="unidentified"
/mol_type="unassigned DNA"
/db_xref="taxon:32644"

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AX022187
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Sequence 8 from Patent WO0046601.
AX033376
AX033376.1 GI:10280150
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   AX022186.1 GI:10045854
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AX033376
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Cech,T.R., Lingner,J., Nakamura,T., Chapman,K.B., Morin,G.B.,
Harley,C.B. and Andrews,W.H.
Human telomerase catalytic subunit
Patent: EP 1333094-A 599 06-AUG-2003;
Geron Corporation (US); University Technology Corporation (US)
Location/Qualifiers
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Human telomerase catalytic subunit
Patent: BF 1333094-A 600 6-AUG-2003;
Geron Corporation (US); University Technology Corporation (US)
Location/Qualifiers
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1 (bases I to 26)
Sechi.T.s., Lingner.J., Nakamura,T., Chapman,K.B., Mori,G.B., Harley,C.B. and Andrews,W.H.
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56;
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Patent: JP 2001081042-A 253 27-MAR-2001;
GERON CORP,UNIVERSITY TECHNOLOGY CORP
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1 BD011296
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Sequence 600 from Patent EP1333094.
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                                                                                            PAT 16-JUL-2002
                                                                                                                                                                                                                                                                                  Kopreski,M.S. and Gocke,C.D.
Method for detection of htr and htert telomerase-associated rna in plasma or serum
Patent: WO 0218652-A 4 07-MAR-2002;
Oncomedx, Inc. (US)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
                                                                                                                                                                                                                         Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Primates, Catarrhini, Hominidae, Homo.
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Method for detection of htr and htert telomerase-associated rna plasma or serum
Patent: WO 0218652-A 5 07-MAR-2002;
Oncomedx, Inc. (US)
Location/Qualifiers
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Pred. No. 56;
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    .26
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    .26
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                                                                                          AX468454 26 bp Sequence 4 from Patent WO0218652. AX468454
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Sequence 5 from Patent WO0218652.
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 1 GAAGGCGTAGGCGCCGTGCTTTTGC 26
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Homo sapiens
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LOCUS DEFINITION ACCESSION VERSION

RESULT 100 AX468455/c

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SOURCE ORGANISM

KEYWORDS

AUTHORS TITLE

JOURNAL

FEATURES

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SOURCE ORGANISM

REFERENCE AUTHORS TITLE

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BD131325
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C07K5/10,
                                                                                                                                                                          A61K38/00, A61K31/7088, A61K39/00, A61K48/00, A61P35/00, A61P43/00,
                                                                                                                                                                                            C07KS/10,
C07KS/107,C07KS/117,C07K7/06,C07K7/08,C07K16/40,C12N9/12, PC
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1 (bases i to 26)
Sechi,T.R., Lingner,J., Nakamura,T., Chapman,K.B., Mori,G.B.,
Harley,C.B. and Andrews,W.H.
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PC C12Q1/02, C12Q1/48, C12Q1/68, G01N33/15, G01N33/50, G01N33/53
G01N33/53,
PC G01N33/566, G01N33/573//C12P21/08, A61K37/02, C12N15/00 CC
SLrandedness: Single;
CC Topology: Linear;
                                                       08/844419
                                                                                                                                                                                                                                                                                     G01N33/566,G01N33/573//C12P21/08,A61K37/02,C12N15/00 CC
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25-APR-1997 US 08/846017,06-MAX-1997 US 08/851843 PR
09-MAX-1997 US 08/854050,14-AUG-1997 US 08/911312 PR
14-AUG-1997 US 08/912951,14-AUG-1997 US 08/915503 PI
R SECHI,JOACHIM LINGNER,TORU NAKAMURA,KAREN B CHAPWAN, PI
PN JP 2001081042-A/253
PP 27-MAR-2001
PP 27-JUL-2000 JP 200227474
PR 01-OCT-1996 US 08/724643,18-APR-1997 US 08/84441
25-APR-1997 US 08/846017,06-MAY-1997 US 08/851843 PR 09-MAY-1997 US 08/851843 PR 09-MAY-1997 US 08/851843 PR 09-MAY-1997 US 08/951312 PR 14-AUG-1997 US 08/951312 PR R SECHI,JOACHIM LINGNER,TORU NAKAMURA,KAREN B CHAPMAN, PI MORIN,
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.larity 100.0%; Pred. No. 56;
Conservative 0; Mismatches 0; Indels
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Patent: JP 2001081042-A 254 27-MAR-2001;
GERON CORP, UNIVERSITY TECHNOLOGY CORP
                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
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                                                                                                                                                             CALVIN B HARLEY, WILLIAM H ANDREWS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human telomerase catalytic subunit.
                                                                                                                                                                                                                                                                                                                                                                                                                              organism="unidentified"
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JP 201081042-A/254
27-MAR-2000
27-JUL-2000 JP 2000227474
01-OCT-1996 US 08/724643
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JP 2001081042-A/254.
                                                                                                                                                    PI CALVIN B HARLEY, WI PC A61K38/00, A61K31/7 PC C07K5/10, C07K5/11 C12N15/09, PC C12Q1/02, C12Q1/48, PC G01N33/53, PC G01N33/53, PC TOPOlogy: Linear; FH KRY Source Lo FT Source 1 FT /organism='Unident
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Matches 26; Conserva
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26 bp DNA linear PAT 18-SEP-2002 Telomerase assay of bodily fluid for cancer screening and evaluation of disease phase and prognosis.
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Method for detecting and inhibiting RNA component of telomerase.
BD023721
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kim, N. W., Wu, F., Kealey, J.T., Pruzan, R. and Weinrich, S.L.
Kim, N.W., Wu, F., Kealey, J.T., Pruzan, R. and Weinrich, S.L.
Method for detecting and inhibiting RNA component of telomerase
Patent: JP 2001507229-A 25 05-JUN-2001;
GERON CORP
PN JP 2001507229-A/25
PN JP 2001507229-A/25
PN JP 1998529003
PR 20-JUN-1997 JP 1998529003
PR 20-DEC-1996 US 08/770564, 20-DEC-1996 US 08/770565 PI
NAM WOO KIM, FRED WU, JAMES T KEALEY, RONALD PRUZAN, SCOTT L PI
                                                                                                                                                                                                                                                    Gaps
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PC C12N5/10.
C C12N5/10.
Strandedness: Single;
C Topology: Linear;
C Topology: Linear;
FH Key Location/Qualifiers.
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56;
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Topology: Linear;
/note= 'hTR forward primer'
Location/Qualifiers.
Location/Qualifiers
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synthetic construct
other sequences; artificial sequences.
1 (bases 1 to 26)
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Best Local Similarity 100.0%; Pred. No. 56;
Matches 26; Conservative 0; Mismatches
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100.0%; Pred. No. 56;
ative 0; Mismatches
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                        source 1. .26 /organism='Unidentified'.
                                                                   Location/Qualifiers
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JP 2002503480-A/3.
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JP 2001507229-A/25.
unidentified
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US09436060A-16.rge.sl

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Topology: Linear;
Modulation of mammalian telomerase by peptide nucleic acids FH
Location/Qualifiers
                                                                                                                                                      PAT 07-SEP-2000
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Modulation of mammalian telomerase by peptide nucleic acids.
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09-APR-1996 US 08/630019
JERRY W SHAY, WOODRING E WRIGHT, MIECZYSLAW A PIATYSZEK, DAVID
                                                                                                                                                                                                                                                                                                                                            Promoter regions of the mouse and human telomerase rna component
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1 (bases 1 to 25)
Shay,J.W., Wright,W.E., Piatyszek,M.A., Corey,D. and Norton,J.C. Modulation of mammalian telomerase by peptide nucleic acids
Patent: JP 2001517929-A 39 09-OCT-2001;
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                                                                                                                                                                                                                                                                                                                                                                           Patent: WO 9938964-A 20 05-AUG-1999;
KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
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63;
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Location/Qualifiers
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                                                                                                                                                      AX019566 25 bp
Sequence 20 from Patent WO9938964.
AX019566
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Strandedness: Single,
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                           175 AATGICAGCIGCIGGCCCGTICGCC 199
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JP 2001517929-A/39
09-OCT-2001
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JP 2001517929-A/39.
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Strovel,J.W., Stamberg,J., Highsmith,E. and Abruzzo,L.V.
Telomerase assay of bodily fluid for cancer screening and
evaluation of disease phase and prognosis
batent: JP 2002503480-A 3 05-FEB-2002;
UNIVERSITY OF MARYLAND BALTIMORE
OS Artificial Sequence
N JP 2002503480-A/3
PD 05-FEB-2002
PF 16-FEB-1999 JP 2000531587
PR 16-FEB-1999 US 60/074793
PR 16-FEB-1999 US 60/074793
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CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
OS Artificial Sequence
N JP 2002509699-A/20
PD 02-ARR-2002
PP 29-JAN-1999 JP 2000529424
PR 29-JAN-1999 GB 9801902.9
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Promoter region of mouse and human telomerase RNA component genes.
BD225817
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Description of Artificial Sequence: F3b, synthesized, Gibco-
BRL
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PC A61P35/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02 PC
.C12Q1/68//C12N9/12,
PC (A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00 CC
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Location/Qualifiers
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    ^coganism="synthetic construct"
    ^mol_type="genomic DNA"
    ^db_xref="taxon:32630"

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    .25
    /organism="synthetic construct"
|mol_type="genomic DNA"
|db_xref="taxon:32630"

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other sequences; artificial sequences.
1 (bases 1 to 25)
Keith, W.N.
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No Unidentified

Catanalogo: Catanalogo: Catanalogo: Catanalogo: Catanalogo: Catanalogo: Linear;

CC Mammalian telomerase

FH Key

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Unclassified.
1 (bases 1 to 28)
1 (lases 1 to 28)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 6548298-A 29 15-APR-2003;
Location/Qualifiers
              unidentified
unclassified.
1 (Dassa 1 to 28)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: 20022272489-A 29 24-SEP-2002;
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Seguence 29 from patent US 6548298.
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Rogeuence 2 from patent US 5846723.

AR063826.

AR063826.1 GI:5993134

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1 (bases 1 to 28)
1 villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Assays for the DNA component of human telomerase Patent: US 5776679-A 28 07-UUL-1998;
Location/Qualifiers
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Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5958680-A 29 28-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5.5%; Score 25; DB 1; Length 28; llarity 100.0%; Pred. No. 71; Conservative 0; Mismatches 0; Indels
               DB 1; Length 25;
63;
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/organism="unknown"
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BD176170
BD176170.1 GI:29121876
JP 2002272489-A/29.
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Promoter region of mouse and human telomerase RNA component genes Patent: JP 200259699-A 13 02-APR-2002;
CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
CANCER PATE CAUGE

BY D 202509699-A/13

BD 02-APR-2002

PF 29-JAN-1998 GB 9801902.9

PI WILLIAM NICOL KEITH
PC CI2NIS/09,A61K31/7105,A61K31/711,A61K35/76,A61K38/00,A61K45/00, PC
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                                                      Promoter region of mouse and human telomerase RNA component genes Patent: JP 2002509699-A 3 02-ARR-2002;
CANCER RESPARCH CAMPAIGN TECHNOLOGY LTD
OS Artificial Sequence
PN JP 2002509699-A/3
PD 02-ARPR-2002
PP 29-JAN-1999 JP 2000529424
PR 29-JAN-1999 GB 9801902.9
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human telomerase RNA component genes.
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C A61P55/00, C12N1/15, C12N1/19, C12N1/21, C12N5/10, C12P21/02
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PC (A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00 CC
Description of Artificial Sequence: Primer
FH Key Location/Qualifiers
1. 24
Creation/Qualifiers
FT source 1. 24
/organism='Artificial Sequence'.
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    .24
    'organism="synthetic construct"
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    'db_xref="taxon:32630"

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other sequences; artificial sequences.
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   (bases 1 to 24)
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Method for detecting and inhibiting RNA component of telomerase.
BD023698
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unclassified.
1 (bases 1 to 30)
Kim,N.W., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.
Method for detecting and inhibiting RNA component of telomerase
Patent: JP 2001507229-A 2 05-JUN-2001;
GERON CORP
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                                                      1 (bases 1 to 30)
Kim,N.Woo., Wu.F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.
Methods for detecting the RNA component of telomerase
Patent: US 5846723-A 2 08-DEC-1998;
Location/Qualifiers
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PD 05-JUN-2001
PP 19-DEC-1997 JP 1998529003
PR 20-DEC-1996 US 08/770564, 20-DEC-1996 US 08/7705
NAM WOO KIM, FRED WU, JAMES T KEALEY, RONALD PRUZAN, SCOTT L
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PC C12N15/09, A61K9/08, A61K31/7105, A61K45/00, A61K48/C
PC C12N5/10, C1201/68, C12O1/68, C12N15/00, C12N5/00 CC
Strandedness: Single;
CC Topology: Linear;
CC Tropology: Linear;
PK Key
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JP 2001507229-A/2.
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Matches 25; Conservative
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DEFINITION ACCESSION RESULT 115 BD023698/c LOCUS

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SOURCE ORGANISM

REFERENCE AUTHORS JOURNAL

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RESULT 116 BD225800/c DEFINITION

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PAT 17-JUL-2003

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Length 24;

PAT 20-DEC-2002

linear

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Promoter regions of the mouse and human telomerase rna component
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Silverman, R.H., Kondo, S., Cowell, J.K., Li, G. and Torrence, P.F.
RNase L activators and antisense oligonucleotides effective to
treat telomerase-expressing malignancies
Patent: US 6468983 A 5 22-OCT-2002;
Location/Qualifiers
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Patent: WO 9938964-A 3 05-AUG-1999;
KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
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100.0%; Pred. No. 71;
ative 0; Mismatches
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Promoter region of mouse and human telomerase RNA component genes. BD25816.1 GI:3303586
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CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
OS Artificial Sequence
N JP 2002509699-A/19
PD 02-APR-2002
PP 29-JAN-1999 JP 2000529424
PR 29-JAN-1999 GB 9801902.9
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Silverman, R.H., Kondo, S., Cowell, J.K., Li, G. and Torrence, P.F.
RNase L activators and antisense oligonucleotides effective to
treat telomerase-expressing malignancies
Patent: US 6469983-A 4 22-OCT-2002;
Location/Qualifiers
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PC (A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00
Description of Artificial Sequence: Primer
FH Key Location/Qualifiers
FT source 1. .24
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5.3%; Score 24; DB 1; Length 24;
Larity 100.0%; Pred. No. 71;
Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                     other sequences; artificial sequences.
1 (bases 1 to 24)
Keith, W.N.
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Sequence 4 from patent US 6468983.
AR241177
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29-JAN-1998 GB 9801902.9
WILLIAM NICOL KEITH
                                                                     46 CTAACCCTAACTGAGAAGGGCGTA 69
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                                                                                       24 CTAACCCTAACTGAGAAGGGCGTA 1
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Matches 24; Conserv
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PAT 07-SEP-2000

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US09436060A-16.rge.sl

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Modulation of mammalian telomerase by peptide nucleic acids.
BD071058
BD071
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In Chases 1 to 24)

(Costable of the control of mammalian telomerase by peptide nucleic acids shay, J.W., wright, W.E., Piatyszek, M.A., Corey, D. and Norton, J.C. Modulation of mammalian telomerase by peptide nucleic acids patent: JP 2001517929-A 24 09-OCT-2001;

GENON CORP OS Unidentified

PN JP 2001517929-A/24

PD 09-OCT-2001

PF 09-APR-1997 JP 1997536487

PR 09-APR-1997 JP 1997536487

COREY,

PI JAMES C NORTON

PC CONTEXT, WASHIX8/16, C1201/68

CC Topology: Linear;

CC Topology: Linear;
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linkages are replaced by N-(2-aminoethyl)glycine units linked
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JP 2001517929-A/24
09-OCT-2001
09-APR-1994 JP 1997536487
09-APR-1996 US 08/630019
JERRY W SHAY, WOODRING E WRIGHT, MIECZYSLAW A PIATYSZEK, DAVID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Topology: Linear; //desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC
                                  Au, J.L. and Wientjes, G.
Compositions active in telomere damage comprising a taxane and
telomerase inhibitor
Patent: WO 0074667-A 5 14-DEC-2000;
Au, Jessie L.S. (US); Wientjes, Guillaume (US)
Location/Qualifiers
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CC linkages are replaced by N-(2-aminoethyl)glycin

CC c nucleotide bases via glycine amino N through a

methylenecarbonyl linker'

FH Key Location/Qualifiers

FT source 1.24

/organism='Unidentified'.
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Location/Qualifiers
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                Patent: WO 9938964-A 13 05-AUG-1999;
KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH
Location/Qualifiers
                               linear
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/db xref="taxon:32630"
/note="primer"
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/organism="synthetic construct"
/mol type="unassigned DNA"
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other sequences; artificial sequences.
                   Sequence 13 from Patent W09938964. AX019559
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Sequence 19 from Patent W09938964.
AX019565
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Sequence 5 from Patent WO0074667,
AX058270
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5.3%; Score 24; DB 1; Length 24; llarity 100.0%; Pred. No. 71; Conservative 0; Mismatches 0; Indels
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BD131326 25 bp DNA linear PAT 18-SEP-2002 Telomerase assay of bodily fluid for cancer screening and evaluation of disease phase and prognosis.
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Description of Artificial Sequence: R3c, synthesized, G1bco-
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JEFFREY W STROVEL, JUDITH STAMBERG, EDWARD HIGHSMITH, LYNNE V
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JP 2002503480-A/4.

Synthetic construct
SM synthetic construct
cother sequences, artificial sequences.

Chasses 1 to 25)
S Strowel, J.W., Stamberg, J., Highsmith, E. and Abruzzo, L.V.
Telomerase assay of bodily fluid for cancer screening and evaluation of disease phase and prognosis
LNIVERSITY OF MARYLAND BALTIMORE
OS Artificial Sequence
PN JP 2002503480-A/4
PD 05-FEB-2002
PP 16-FEB-1999 US 60/074793
PR 16-FEB-1999 US 60/074793
PI JEFREY W STROVEL, JUDITH STAMBERG, EDWARD HIGHSMITH, LYNNE
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/organism='Artificial Sequence'
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                                                                                        DB 1;
71;
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                                                                                        5.3%; Score 24; DB 100.0%; Pred. No. 71; iive 0; Mismatches
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JP 2001517929-A/25.
                                                                                                                            Conservative
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                  PAT 27-AUG-2002
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Synthetic construct

Synthetic construct

other sequences; artificial sequences.

E 1 (bases 1 to 24)

S Silverman, R. H., Kondo, S., Cowell, J. K., Li, G. and Torrence, P. F.

RNase L activators and antisense oligonucleotides effective to treat telomerses-expressing malignancies

L Patent: JP 2001524100-A 5 27-NOV-2001;

THE CLEVELAND CLINIC FOUNDATION, NATIONAL INSTITUTES OF HEALTH OS Artificial Sequence

PN JP 2001524100-A/5
                                                                                                                                                                                                                                                                                      OS Artificial Sequence
PN JP 2001524100-A/4
PD 27-NOV-2001
PF 13-APR-1999 US 60/044507,03-FEB-1998 US 09/018125 PI
ROBERT H SILVERMAN, SEIJI KONDO, JOHN K COWELL, GUIYING LI, PAUL F
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PN JP 2001524100-A/5
PD 27-NOV-2001
PF 13-APR-1998 JP 1998546125
PR 21-APR-1997 US 60/044507,03-FEB-1998 US 09/018125 PI
ROBERT H SILVERMAN, SEIJI KONDO, JOHN K COWELL, GUIYING LI, PAUL F
                                                                                                                    synthetic construct
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other sequences; artificial sequences.

1 (bases 1 to 24)
Silverman, R.H., Kondo, S., Cowell, J.K., Li, G. and Torrence, P.F.
RNase L activators and antieense oligonucleotides effective to
treat telomerase-expressing malignancies
Patent: JP 2001524100-A 4 27-NOV-2001;
THE CLEVELAND CLINIC FOUNDATION, NATIONAL INSTITUTES OF HEALTH
DN JP 2001524100-A/4
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C07H21/00,C07H21/02,C12Q1/68,A61K48/00
Description of Artificial Sequence: primer
Key
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Description of Artificial Sequence: primer
Key

    .24
    Organism="synthetic construct"
|mol_type="genomic DNA"
|db_xref="taxon:32630"

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ilarity 100.0%; Pred. No. 71;
Conservative 0; Mismatches
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JP 2001524100-A/4.
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BD084641/c
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH
Location/Qualifiers
               Length 25;
                                                                 Indels
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4.9%; Score 22; DB 1; Length 22;
Best Local Similarity 100.0%; Pred. No. 90;
Matches 22; Conservative 0; Mismatches 0; Indels
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/organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/note="Oligonucleotide"
               DB 1;
87;
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Villeponteau, B., Feng, J., Funk, W. and Assays for the DNA component of human Patent: US 5776679-A 32 07-UUL-1998;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                             other sequences; artificial sequences.
             5.1%; Score 23; DB 100.0%; Pred. No. 87; Live 0; Mismatches
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LOCUS AR059220
DEFINITION Sequence 27 from patent US 5837857.
                                                                                                                                                                                                                                                                                 Sequence 44 from Patent W09938964.
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                                                                                                             1 GGGTTGCGGAGGGTGGGCCTGGG 23
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/organism="unknown"
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     Query Match
Best Local Similarity 100.0
Matches 23; Conservative
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ACCESSION
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FEATURES
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AUTHORS
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Promoter region of mouse and human telomerase RNA component genes.
BD225841
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                                                                                                                                                                                                                                                                         Topology: Linear; / desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC / desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC / desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC / desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC / desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC / desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC / desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC / desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC / desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC / desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC / desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC / desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC / desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC / desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC / desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC / desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC / deoxy(ribose- CC / deoxy(r
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PD 02-APR-2002
                                                                                                09-APR-1997 JP 1997536487
09-APR-1996 US 08/630019
JERRY W SHAY, WOODRING E WRIGHT, MIECZYSLAW A PIATYSZEK, DAVID
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AGIK48/00,

PC AGIP35/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,L1622.

C12Q1/67(C12N9/12,

C1Q1/64(X15/7)C12N1/12,C12N15/00,A61K37/02,C12N5/00

Description of Artificial Sequence:Oligonucleotide FH K
Location/Qualifiers

Location/Qualifiers
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                        C linkages are replaced by N-(2-aminoethyl)glycin CC to to to CC nucleotide bases via glycine amino N through a methylenecarbonyl linker'
FH Key Location/Qualifiers
FT source 1. .23
FT /organism='Unidentified'.
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80;
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1 (bases 1 to 25)
Keith, W.N.
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JAMES C NORTON
CO7K14/00,A61K38/16,C12Q1/68
Strandedness: Single;
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/organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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29-JAN-1998 GB 9801902
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JP 2002509699-A/44.
synthetic construct
synthetic construct
                          Unidentified
JP 2001517929-A/25
09-OCT-2001
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Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5958680-A 42 28-SEP-1999;
Location/Qualifiers
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Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5958680-A 43 28-SEP-1999;
Location/Qualifiers
                                                                                                                     linear
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4.9%; Score 22; DB 1;
Best Local Similarity 100.0%; Pred. No. 90;
Matches 22; Conservative 0; Mismatches (
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Sequence 42 from patent US 5958680.
AR075545.
AR075545.1 GI:10002291
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4.9%; Score 22; DB
Best Local Similarity 100.0%; Pred. No. 90;
Matches 22; Conservative 0; Mismatches
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Sequence 43 from patent US 5958680.
AR075546
AR075546.1 GI:10002292
    Mismatches
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                           CCCTCC 204
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Mammalian telomerase.
BD176150
BD176150.1 GI:29121854
JP 2002272489-A/9.
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unclassified.
    Conservative
                           183 CTGCTGGCCCGTTCG
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AR075545/c
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AR075546/c
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1 (bases 1 to 22)
11leponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5837857-A 28 17-NOV-1998;
Location/Qualifiers
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Unclassified.
Unclassified.
1 (bases 1 to 22)
111eponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5958680-A 37 28-SEP-1999;
Location/Qualifiers
                                                    Unknown.
Unclassified.
1 (bases 1 to 22)
Villeponteau,B., Feng,J., Funk,W. and Andrews,W.H. Mammalian telomerase
Patent: US 5837857-A 27 17-NOV-1998;
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4.9%; Score 22; DB 1;
Best Local Similarity 100.0%; Pred. No. 90;
Matches 22; Conservative 0; Mismatches
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4.9%; Score 22; DB 1;
Best Local Similarity 100.0%; Pred. No. 90;
Matches 22; Conservative 0; Mismatches
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Sequence 37 from patent US 5958680.
AR075540
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Sequence 28 from patent US 5837857.
AR059221
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/organism="unknown"
/wol_type="unassigned DNA"
                                                                                                                                   Location/Qualifiers
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               GI:5984797
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  AR059220
AR059220.1
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OS Unidentified
PN JP 200222489-A/32
PD 24-SEP-2002
PP 24-SEP-2002
PP 06-MAR-2002 JP 2002061125
PR 07-JUL-1994 US 08/330123 PR 07-JUL-1994 US 08/330123 PR 07-JUL-1995 US 08/42802,07-JUN-1995 US 08/482115 PI BRYANT
VILLEPONTEAL, JUNII FENG, WALTER FUNK, WILLIAM HANDREMS PC C12N15/09, C12N9/99, C12Q1/68, G01N33/53, G01N33/56, C12N15/00 CC Strandedness: Single;
CC Topology: Linear;
CC Mammalian telomerase
FH Key Location/Qualifiers
FT source / Organism='Unidentified'.
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Unclassified
1 (Dassa 1 to 22)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: 2002272489-A 32 24-SEP-2002;
                                                                                                                                                                                           linear
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Unclassified.
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Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: US 5583016-A 5 10-DEC-1996;
Locarion/Qualifiers
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Pred. No. 90;
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100.0%; Pred. No. 90;
iive 0; Mismatches (
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4.9%; Score 22; DB
Best Local Similarity 100.0%; Pred. No. 90;
Matches 22; Conservative 0; Mismatches
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/organism="unknown"
/mol_type="unassigned DNA"
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Sequence 5 from patent US 5583016.
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/organism="unidentified"
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/db_xref="taxon:32644"
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               54 AACTGAGAAGGGCGTAGGCGCC 75
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JP 2002272489-A/32.
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PN JP 2002272489-A/10
PD 24-SEP-2002
PF 6-MAR-2002 JP 2002061125
PF 07-JUL-1994 US 08/472102,27-OCT-1994 US 08/330123 PR 07-JUL-1995 US 08/472802,07-JUL-1995 US 08/482115 PI BRYANT VILLEPONTEAU, JUNLI FRNG, WALTER FUNK, WILLIAM H ANDREWS PC C12N15/09,C12N9/99,C12Q1/68,G01N33/53,G01N33/566,C12N15/00 CC CTOPOLOGY: Linear; CC Mammalian telomerase FH Key Location/Qualifiers
FT source Location/Qualifiers
FT source /organism='Unidentified'.
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  1 (bases 1 to 22)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: JP 2002272489-A 9 24-SEP-2002;
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4.9%; Score 22; DB 1; Length 22;
Best Local Similarity 100.0%; Pred. No. 90;
Matches 22; Conservative 0; Mismatches 0; Indels
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Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: JP 2002272489-A 10 24-SEP-2002;
GERON CORP
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/organism='Unidentified'
Location/Qualifiers
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JP 2002272489-A/10.
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BD176151/c
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1 (bases 1 to 22)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerae
Patent: US 6548298-A 41 15-APR-2003;
Location/Qualifiers
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Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: US 6548299-A 42 15-APR-2003;
                                                                                                                        Length 22;
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Best Local Similarity 100.0%; Pred. No. >v,
Matches 22; Conservative 0; Mismatches
   and Pruzan,R.A. Methods for purifying telomerase Patent: US 6545113-A 2 08-APR-2003; Location/Qualifiers
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Sequence 42 from patent US 6548298.
AR306490
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4.9%; Score 22; DB
Best Local Similarity 100.0%; Pred. No. 90;
Matches 22; Conservative 0; Mismatches
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/organism="unknown"
/mol_type="unassigned RNA"
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/mol_type="genomic DNA"
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Weinrich,S.L., Atkinson,E.M. III, Lichtsteiner,S.P., Vasserot,A.P. and Pruzan,R.A.
Purified telomerase
Patent: US 6517834-A 2 11-FEB-2003;
Location/Qualifiers
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1 (bases 1 to 22)
Weinrich, S.L., Atkinson, E.M. III, Lichtsteiner, S.P., Vasserot, A.P.
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Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5583016-A 6 10-DEC-1996;
Location/Qualifiers
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Sequence 2 from patent US 6545133.
AR305068
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Sequence 6 from patent US 5583016.
131753.1 GI:1822544
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22 CTAACCCTAACTGAGAAGGGCG 1
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Best Local Similarity 100.0
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SOURCE ORGANISM

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RESULT 143 AR279617/c DEFINITION

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PAT 12-JUN-2003

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RESULT 144 AR305068/c

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OS Unidentified
PN JP 2002272489-A/27
PD 24-SEP-2002
PF 06-MAR-2002 JP 2002061125
PF 06-MAR-2002 JP 2002061125
PF 07-JUL-1994 US 08/472802,07-001-1995 US 08/482115 PI BRYANT
07-JUL-1995 US 08/472802,07-001-1995 US 08/482115 PI BRYANT
VILLEPONTEAL, UNMLI FENG, WALTER FUNK, WILLIAM HANDREWS PC
C12N15/09, C12N9/99, C12Q1/68, G01N33/53, G01N33/566, C12N15/00 CC
Strandedness: Single,
CC Mammalian telomerase
FT source
I.ocation/Qualifiers
FT source
/organism='Unidentified'.
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Unclassfiled.
1 (bases 1 to 21)
Villeponteau.B., Feng,J., Funk,W. and Andrews,W.H.
Mammalian telomerasa.
Patent: US 625635-A 25 10-JUL-2001;
Location/Qualifiers
                              4.7%; Score 21; DB 1; Length 21; 100.0%; Pred. No. 1e+02;
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llarity 100.0%; Pred. No. 1e+02;
Conservative 0; Mismatches 0; Indels
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1 (bases 1 to 21)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: JP 2002272489-A 27 24-SEP-2002;
GERON CORP
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                                         larity 100.0%; Pred. No. 1e-
Conservative 0; Mismatches
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Sequence 25 from patent US 6288535.
AR161927
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JP 2002272489-A/27.
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BD176168
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Unclassified.
I (bases It o 21)
I (lasses It o 21)
Assays for the DNA component of human telomerase
Patent: US 5776679-A 25 07-UL-1998;
Location/Qualifiers
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Unclassified.
Unclassified.
1 (bases 1 to 21)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5837857-A 25 17-NOV-1998;
Location/Qualifiers
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4.7%; Score 21; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 21; Conservative 0; Mismatches 0; Indels
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1 (bases 1 to 21)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Patent: US 5958680-A 26 28-SEP-1999;
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100.0%; Pred. No. 10.
... 0; Mismatches
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Sequence 25 from patent US 5837857.
AR059218
                           AR016057 21 bp Sequence 25 from patent US 5776679.
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AR075529
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                                                                      AR016057.1 GI:3972334
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Matches 21; Conservative
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PAT 31-AUG-2000

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PAT 27-AUG-2002

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1 (bases 1 to 22)
Weinrich,S.L., Atkinson,E.M. III, Lichtsteiner,S.P., Vasserot,A.P.,
Pruzan,R.A. and Kealey,J.T.
Purified telomerase
Patent: US 596856-A 3 19-OCT-1999;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD calde; Panitooideae; Andropogoneae; Zea.

1 (bases 1 to 22)
Weinrich,S.L., Iii,E.M.A., Lichtsteiner,S.P., Vasserot,A.P., Pruzan,R.A. and Kealey,J.T.
Pruzan,R.A. and Kealey,J.T.
Parent: JP 2001509681-A 3 24-JUL-2001;
GERON CORP
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unidentified
unidentified
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1 (bases 1 to 22)
Atkinson, E.M. and Kealey, J.T.
PURITED TELOMERASE
PARENT: WO 9845450-A 3 15-OCT-1998;
GERON CORP (US)
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Sequence 3 from patent US 5968506.
AR079890
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JP 2001509681-A/3.
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Purified telomerase.
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BD058134/c
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Unclassified.
Unclassified.
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 6548298-A 25 15-APR-2003;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4.7%; Score 21; DB 1; Length 21; ilarity 100.0%; Pred. No. 1e+02; Conservative 0; Mismatches 0; Indels
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Villeponteau B., Feng,J., Funk,W. and Andrews,W.H.
Mammalian telomerase
Patent: US 5583016-A 25_10-DEC-1996;
                   Length 21;
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                Query Match 4.7%; Score 21; DB 1; Best Local Similarity 100.0%; Pred. No. 1e+02; Matches 21; Conservative 0; Mismatches 0
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Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 21; Conservative 0; Mismatches 0
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131772
131772.1 GI:1822563
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Sequence 25 from patent US 6548298.
AR306475.1 GI:31696314
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Sequence 3 from Patent WO9845450.
A84593.1 GI:6733509
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131772/c
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AR306475/c
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PAT 29-SEP-1999
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                                                                                                                                                                                                                                                                                                                                       Unclassified.
1 (bases 1 to 20)
Villeponteau.B., Feng,J., Funk,W. and Andrews,W.H.
Mammalian telomerase
Patent: US 5837857-A 7 17-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     [bases 1 to 20)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5837857-A 26 17-NOV-1998;
Location/Qualifiers
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                                                           Length 20;
                                                                                                                                                                                                                                    linear
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4.4%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 1.1e+02;
Matches 20; Conservative 0; Mismatches 0; Indels
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                                                           Query Match
4.4%; Score 20; DB 1; Lv
Best Local Similarity 100.0%; Pred. No. 1.1e+02;
Matches 20; Conservative 0; Mismatches 0;
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Sequence 29 from patent US 5837857.
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Sequence 26 from patent US 5837857.
AR059219.1 GI:5984796
                                                                                                                                                                                                                     AROS9200 20 bp
Sequence 7 from patent US 5837857.
AROS9200
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            41 TTTGTCTAACCCTAACTGAG 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 GGTTGCGGAGGGTGGGCCTG 21
                                                                                                                         2 GGTTGCGGAGGGTGGGCCTG 21
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Unclassified.
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AR059219/c
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AR059200/c
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PN JP 2001509681-A/3
PD 24-JUL-2001
PF 04-ARP.1997 JP 1998542718
PI SCOTT L WEINRICH. EDWARD M ATKINSON III.SERGE P LICHTSTEINER,
PI ALAIN P VASSEROT, RONALD A PRUZAN, JAMES T KEALEY PC
C12N15/54, C12N9/12, C07K16/40, C12Q1/68, C07K14/47 CC Strandedness:
Single;
CC Topology: Linear;
CC /mod_base= OTHER
CC /mod_base= OTHER
CC /note= 'N = biotinylated C'
CC /note= 'N = biotinylated C'
CC /note= 'Oligonucleotide P3'
FH Key Location/Qualifiers
FT modified base 1:
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1 (bases 1 to 20)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Assays for the DNA component of human telomerase Patent: US 5776679-A 7 07-UUL-1998;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                   Query Match 4.7%; Score 21; DB 1; Length 22; Best Local Similarity 100.0%; Pred. No. 1e+02; Matches 21; Conservative 0; Mismatches 0; Indels
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100.0%; Pred. No. 1.1e+02;
tive 0; Mismatches 0;
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Atkinson, E.M. and Kealey, J.T.
PURIFIED FILCHMERASE
PATENT: WO 9845450-A 14 15-OCT-1998;
GERON CORP (US)

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Sequence 7 from patent US 5776679.
AR016039.1 GI:3972316
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Sequence 14 from Patent WO9845450.
A84604
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                                                                                                                                                                                                                                                                                                                                                                   46 CTAACCCTAACTGAGAAGGGC 66
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AR016039/c
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PAT 29-SEP-1999

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1 (bases 1 to 20)
Kim, N.Woo., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.
Kim,N.Woo., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.
Methods for detecting the RNA component of telomerase
Patent: US 5846723-A 13 08-DEC-1998;
Location/Qualifiers
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Kim,N.Woo., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.

Methods for detecting the RNA component of telomerase

Patent: US 5846723-A 7 08-DEC-1998;

Location/Qualifiers
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100.0%; Pred. No. 1.1e+02;
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4.4%; Score 20; DB 1; L
Best Local Similarity 100.0%; Pred. No. 1.1e+02;
Matches 20; Conservative 0; Mismatches 0;
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 0; Mismatches
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                                                                                                                      20 bp
Sequence 7 from patent US 5846723.
AR063831.1 GI:5993139
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AR063837
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Sequence 8 from patent US 5958680.
AR075511
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                             GAAGAGTTGGGCTCTGTCAG 319
                                                                                                                                                                                                                                                                                                                                                                                                                                 290 CTGCCACCGCGAAGAGTTGG 309
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                                                      GAAGAGTTGGGCTCTGTCAG 1
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AR063831/c
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AR075511/c
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AR063837/c
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Unclassified.
Unclassified.
I (bases I to 20)
Kim,N. Woo., Wu.F., Kealey,J.T., Fruzan,R. and Weinrich,S.L.
Methods for detecting the RNA component of telomerase
Patent: US 5846723-A 3 08-DEC-1998;
Location/Qualifiers
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Kim,N.Woo., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.

Methods for detecting the RNA component of telomerase

Patent: US 5846723-A 6 08-DEC-1998;

Location/Qualifiers
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                                                                                1 (bases 1 to 20)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: US 5837857-A 29 17-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4.4%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 1.1e+02; tive 0; Mismatches 0; Indels
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Sequence 3 from patent US 5846723.
AR063827.1 GI:5993135
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20 bp
AR063830 GIrom patent US 5846723.
AR063830.1 GI:5993138
                                                                                                                                          Location/Qualifiers
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              GI:5984799
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Best Local Similarity
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Unclassified.
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Matches 20; Conserva
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AR059222.1
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AR063827/c
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AR063830/c
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PAT 17-OCT-2001
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                                                                                                                                                                     Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: US 625835-A 7 10-UUL-2001;
Location/Qualifiers
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                                            20 bp
Sequence 7 from patent US 6258535.
AR161909
AR161909.1 GI:16228923
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="unidentified"
/mol_type="genomic DNA"
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JP 2002272489-A/8.
unidentified
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Best Local Similarity 100..
These 20; Conservative
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Matches 20; Conservative
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Unclassified.
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                         RESULT 169
AR161909/c
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BD176149/c
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Weinrich, S.L., Atkinson, B.M. III, Lichtsteiner, S.P., Vasserot, A.P.,
Pruzan, R.A. and Kaaley, J.T.
Purified telomerase
Parent: US 5968506-A 7 19-OCT-1999;
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1 (bases 1 to 20)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: US 5958680-A 8 28-SEP-1999,
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                    Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H., Willeponteau, B., Mammalian telomerase
Patent: US 5958680-A 41 28-SEP-1999;
Location/Qualifiers
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4.4%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 1.1e+02;
Matches 20; Conservative 0; Mismatches 0; Indels
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Best Local Similarity 100.0%; Pred. No. 1.1e+02;
Matches 20; Conservative 0; Mismatches 0;
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/organism="unknown"
/mol_type="unassigned DNA"
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Sequence 7 from patent US 5968506.
AR079894

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1 (bases 1 to 20)
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Unclassified.
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PF 29-JAN-1999 JP 2000529424
PR 29-JAN-1998 GB 9801902.9
PI WILLIAM NICOL KEITH
PC C12N15/09,A61K31/7105,A61K31/711,A61K35/76,A61K38/00,A61K45/00,PC
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Promoter region of mouse and human telomerase RNA component genes.
BD225842
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CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
OS ARLifficial Sequence
PN JP 2002509699-A/45
PP 02-APR-2002
PP 29-JAN-1999 JP 2000529424
PR 29-JAN-1999 GB 9801902.9
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PC A61P35/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02 PC
C12Q1/68//C12N9/12,
PC (A61X35/76,A61X31:522),C12N15/00,A61X37/02,C12N5/00 CC
Description of Artificial Sequence:Oligonucleotide FH Key
Location/Qualifiers
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            ce 1. .20 /organism='Artificial Sequence'
Location/Qualifiers
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100.0%; Pred. No. 1.1e+02;
rative 0; Mismatches 0; Indels
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    .20
    ^coganism="synthetic construct"
    ^mol_type="genomic DNA"
    ^db_xref="taxon:32630"

Location/Qualifiers

    .20
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synthetic construct
other sequences; artificial sequences.
1 (bases 1 to 20)
Keith, W.N.
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Sequence 4 from patent US 5583016.
131751
131751.1 GI:1822542
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JP 2002509699-A/45.
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BD225842
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PN JP 2002272469-A/11
PD 24-SEP-2002 JP 2002061125
PF 06-MAR-2002 JP 2002061125
PR 07-JUJ-1994 US 08/72102, 27-OCT-1994 US 08/330123 PR 07-JUJ-1995 US 08/482115 PI BRYANT VILLEPONTEAU, JUNIA FENG, WALTER FUNK, WILLIAM H ANDREWS PC CIANIS/09, C12N9/99, C12Q1/68, G01N33/53, G01N33/566, C12N15/00 CC Strandedness: Single;
CC Topology: Linear;
CC Mammalian telomerase
FH Key Location/Qualifiers
FT source /organism='Unidentified',
                                               PAT 18-MAR-2003
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human telomerase RNA component genes.
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PC (A61K35/76,A61K1.522).C12N15/00,A61K37/02,C12N5/00 CC

Description of Artificial Sequence: Primer
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Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Mammalian telomerase
Patent: JP 2002272489-A 11 24-SEP-2002;
GERON CORP
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Location/Qualifiers
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ilarity 100.0%; Pred. No. 1.1e+02;
Conservative 0; Mismatches 0;
                                               DNA
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JP 2002509699-A/15.
synthetic construct
synthetic construct
other sequences; artificial sequences.

    .20
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/db_xref="taxon:32644"

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                                               20 bp
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                                                              Mammalian telomerase.
BD176152
BD176152.1 GI:29121856
JP 2002272489-A/11.
unidentified
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                                               BD176152
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              RESULT 171
BD176152/c
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PAT 12-JUN-2003
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Patent: WO 9938964-A 15 05-AUG-1999;
KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
                                                                                                                                                    Unkrown.
Unclassfiled.
1 (bases 1 to 20)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H.
Wammalian telomerash
Mammalian telomerash
Patent: US 6548298-A 40 15-APR-2003;
Locaion/Qualifiers
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4.4%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 1.1e+02;
Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                           Length 20;
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ilarity 100.0%; Pred. No. 1.1e+02;
Conservative 0; Mismatches 0;
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/noTe="primer"
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synthetic construct
other sequences; artificial sequences.
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                                                                                                                                                                                                                                                                 /organism="unknown"
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Sequence 45 from Patent WO9938964.
AX019591.1 GI:10043505
                                                         AR306488 20 bp Sequence 40 from patent US 6548298.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AX019561 20 bp
Sequence 15 from Patent W09938964.
AX019561 GI:10043475
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                                                                                                    AR306488.1 GI:31696327
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synthetic construct
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Matches 20; Conserv
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Unknown.
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                              RESULT 177
AR306488/c
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AX019591
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Unclassified.

1 (bases 1 to 20)
Willeponteau B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: US 5583016-A 4 10-DEC-1996;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                    Unclassified.

1 (Dases 1 to 20)
Villeponteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: US 5583016-A 7 10-DEC-1996;
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Unclassified.
1 (bases 1 to 20)
1 (baleporteau, B., Feng, J., Funk, W. and Andrews, W.H. Mammalian telomerase
Patent: US 6548298-A 7 15-APR-2003;
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Best Local Similarity 100.0%; Pred. No. 1.1e+02;
Matches 20; Conservative 0; Mismatches 0;
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4.4%; Score 20; DB 1; L.
Best Local Similarity 100.0%; Pred. No. 1.1e+02;
Matches 20; Conservative 0; Mismatches 0;
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/mol_type="unassigned DNA"
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Sequence 7 from patent US 6548298.
AR306459
AR306459.1 GI:31696298
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Sequence 7 from patent US 5583016.
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/organism="unknown"
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AR306459/c
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FEATURES
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131754/c
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20 bp DNA linear PAT 27-AUG-2002
Method for detecting and inhibiting RNA component of telomerase.
BD023703
BP023703.1 GI:22564926
JP 2001507229-A/7.
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WEINRICH
PC C12N15/09, A61K9/08, A61K31/7105, A61K45/00, A61K48/00, A61P35/00,
PC C12N15/10,
PC C12N9/12, C12Q1/68, C12Q1/68, C12N15/00, C12N5/00 CC
Strandedness: Single;
CC Topology: Linear;
CC Topology: Linear;
PC Note= 'oligo 14ab'
PH Key
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C12NS/10,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         unidentified unclassified.

unclassified.
1 (bases 1 to 20)
Kim, W., W., F., Kealey, J.T., Pruzan, R. and Weinrich, S.L.
Kim, More detecting and inhibiting RNA component of telomerase Patent: JP 2001507229-A 6 05-JUN-2001;
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PD 05-JUN-2001
PF 19-DEC-1999 JJP 1998529003
PR 20-DEC-1996 US 08/770565 PI
NAM WOO KIM, FRED WU, JAMES T KEALEY, RONALD PRUZAN, SCOTT L PI
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PC C12N15/09, A61K9/08, A61K31/7105, A61K45/00, A61K4)
PC C12N5/10,
PC C12N9/12, C12Q1/68, C12Q1/68, C12N15/00, C12N5/00 (Strandedness: Single;
CC Topology: Linear;
CC Topology: Linear;
CC Topology: Linear;
PF Key Location/Qualifiers.
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4.4%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 1.1e+02;
Matches 20; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                  4.4%; Score 20; DB 1; Le
ilarity 100.0%; Pred. No. 1.1e+02;
Conservative 0; Mismatches 0;
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/mol_type="genomic DNA"
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JP 2001507229-A/6.
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Method for detecting and inhibiting RNA component of telomerase.
BD021699
   Keith, W.N. Promoter regions of the mouse and human telomerase rna component
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Kim,N.W., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.
Method for detecting and inhibiting RNA component of telomerase
Patent: JP 2001507229-A 3 05-JUN-2001;
GERON CORP
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PD 05-JUN-2001
PF 19-DEC-19997 JP 1998529003
PR 20-DEC-1996 US 08/770565 PI
NAM WOO KIM, FRED WU, JAMES T KEALEY, RONALD PRUZAN, SCOTT L PI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Au, J.L. and Wientjes, G.
Compositions active in telomere damage comprising a taxane and telomerase inhibitor
Patent: WO 0074667-A 6 14-DEC-2000;
Au, Jessie L.S. (US)
Location/Qualifiers
                                                                                                                                                                                                                                                                      Gaps
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                                                    Scales

KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)

Location/Qualifiers

1. 20

/organism="synthetic construct"
/mol_type="unassigned DNA"
/db xref="taxon:32630"
/noTe="oligonucleotide"
                                                                                                                                                                                                                           4.4%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 1.1e+02; ive 0; Mismatches 0; Indels
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4.4%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 1.1e+02;
Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
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/organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/note="primer/probe"
                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  synthetic construct
synthetic construct
other sequences; artificial sequences.
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Sequence 6 from Patent WO0074667.
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JP 2001507229-A/3.
unidentified
unidentified
unclassified.
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AX058271.1 GI:12310770
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Matches 20; Conservative
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BD023699/c
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AX058271
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                                                                                                                                                                                                                                                                                                                                                                       Zea mays
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Eukaryota; Poaceae; PACCAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PN JP 2001509681-A/11
PD 24-JUL-2001
PF 04-APR-1997 1998542718
PF SCHAPR-1997 1 WEINERH SENARD M ATKINSON III, SERGE P LICHTSTEINER, PI ALAIN P VASSEROT, RONALD A PRUZAN, JAMES T KEALEY PC C12N15/54, C12N9/12, C07K16/40, C12Q1/68, C07K14/47 CC Strandedness:
                                                                                                                                                                                                                                                                                                                                                                                                                        Spermatophyta; Magnoliophyta; Lillopsida; Poales; Poaceae; PACCAD clade; Panicoideae; Andropogoneae; Zea.

1 (bases 1 to 20)

Reinrich, S.L., Ili, E.M.A., Lichtsteiner, S.P., Vasserot, A.P., Pruzan, R.A. and Kealey, J.T.

Puried telomerase
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4.4%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 1.1e+02;
Matches 20; Conservative 0; Mismatches 0; Indels
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GERON CORP
                                                                                                                                                                                                                    DNA
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/db_xref="taxon:4577"
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                                159 TCTAGAGCAACAAAAATG 178
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BD058142.1 GI:22603748
JP 2001509681-A/11.
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Purified telomerase.
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BD058142/c
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C12N5/10,
                        unidentified
unclassified.
1 (bases 1 to 20)
Kim,N.W., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.
Method for detecting and inhibiting RNA component of telomerase
Patent: JP 2001507229-A 7 05-JUN-2001;
GERON CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BD023709 20 bp DNA linear PAT 27-AUG-2
Method for detecting and inhibiting RNA component of telomerase.
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1 (bases 1 to 20)

Kim,N.W., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.

Kim,N.W. Getecting and inhibiting RNA component of telomerase Patent: JP 2001507229-A 13 05-JUN-2001;

GERRON CORP

PN JP 2001507229-A/13

PD 05-JUN-2001

PF 19-DEC-1997 JP 1998529003

PR 20-DEC-1996 US 08/770565 PI

NAM WOO KIM,FRED WU,JAMES T KEALEY,RONALD PRUZAN,SCOTT L PI

WEINRICH
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PD 05-UTM-2001
PF 19-DEC-1997 UP 1998529003
PR 20-DEC-1996 US 08/770564, 20-DEC-1996 US 08/770565
NAM WOO KIM, FRED WU, JAMES T KEALEY, RONALD PRUZAN, SCOTT L PI
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PC C12N5/10,
PC C12N9/12,C12Q1/68,C12Q1/68,C12N15/00,C12N5/00 CC
Strandedness: Single;
CC Topology: Linear;
CC Topology: Linear;
FH Key Location/Qualifiers.
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PC C12N5/10.
C C12N5/10.
Strandedness: Single;
CC Topology: Linear;
PH Key Location/Qualifiers.
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4.4%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 1.1e+02;
Matches 20; Conservative 0; Mismatches 0;
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    .20
    /organism="unidentified"
    /mol_type="genomic DNA"
    /db_xref="taxon:32644"

    .20
    /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"

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JP 2001507229-A/13.
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BD023709/c
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76 GIGCTITIGCICCCCCCC 94
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JP 2001507229-A/9.
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BD023705/c
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KEYWORDS
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BD071093
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                                                  linkages are replaced by N-(2-aminoethyl)glycine units linked
             1 (bases 1 to 19)
Silverman, R.H., Kondo, S., Cowell, J.K., Li, G. and Torrence, P.F.
RNase L activators and antisense oligonucleotides effective to
treat telomerase-expressing malignancies
Patent: US 646993-A 2 22-0CT-2002;
Location/Qualifiers
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Kim,N.Woo., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.

Methods for detecting the RNA component of telomerase

Patent: US 5846723.4 9 08-DEC-1998;

Location/Qualifiers
                                             C linkages are replaced by N-(2-aminoethyl)glycincCC to to to to CC nucleotide bases via glycine amino N through a methylenecarbonyl linker'
FH Key Location/Qualifiers 1. 20
FT source /organism='Unidentified'.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4.2%; Score 19; DB 1; Length 19;
100.0%; Pred. No. 1.2e+02;
                                                                                                                                                                                                                         Score 20; DB 1; Length 20;
Pred. No. 1.1e+02;
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100.0%; Pred. No.
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/organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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/organism="unknown"
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Strandedness: Single;
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(NAM WOO KIM, FRED WU, JAMES T KEALEY, RONALD PRUZAN, SCOTT L PI

(NEINSTCH WEINSTCH PC 1208/10, A61K9/08, A61K31/7105, A61K45/00, A61P35/00, PC C12NS/10, C12US/12, C12Q1/68, C12U15/00, C12NS/10, C1
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Method for detecting and inhibiting RNA component of telomerage.
BD023705
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1 (bases 1 to 19)
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1 (hase fire, Kealey,J.T., Pruzan,R. and Weinrich,S.L. Method for detecting and inhibiting RNA component of telomerase Patent: JP 2001507229-A 9 05-JUN-2001;
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    /organism="unidentified"
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JP 2001517929-A/59
09-OCT-2001
09-APR-1997 JP 1997536487
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361 AGGCCGCAGGAAGAGGAAC 379
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Strandedness: Single;
Topology: Linear;
Modulation of mammalian telomerase by peptide nucleic acids FH source
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JERRY W SHAY, WOODRING E WRIGHT, MIECZYSLAW A PIATYSZEK, DAVID
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synthetic construct
construct
other sequences; artificial sequences.

(bases 1 to 19)
Silverman, R.H., Kondo, S., Cowell, J.K., Li, G. and Torrence, P.F.
RNase L activators and antisense oligonucleotides effective to
treat telomerase-expressing malignancies
Patent: JP 2001524100-A 2 27-NOV-2001;
THE CLEVELAND CLINIC FOUNDATION, NATIONAL INSTITUTES OF HEALTH
Dy JP 2001524100-A/2
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PP 27-NOV-2001

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ROBERT H SILVERMAN,SEIJI KONDO,JOHN K COWELL,GUIYING LI,PAUL J

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/organism="synthetic construct"
/mol type="genomic DNA"
/db_kref="taxon:32630"
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/db_xref="taxon:32644"
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JAMES C NORTON
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Weinrich, S.L., Atkinson, E.M. III, Lichtsteiner, S.P., Vasserot, A.P., Pruzan, R.A. and Kealey, J.T.
Purified telomerae Purified telomerae Patent: US 5968506-A 14 19-0CT-1999;
Location/Qualifiers
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Spermatophyta, Magnoliophyta, Liliopsida, Poales, Poaceae, PACCAD
clade, Panicoideae, Andropogoneae, Zea.
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Weinrich, S.L., Iii, E.M.A., Lichtsteiner, S.P., Vasserot, A.P., Pruzan, R.A. and Kealey, J.T.
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Sequence 14 from patent US 5968506.
AR079898
                                                                                                                  Patent: WO 9845450-A 7 15-OCT-1998;
GERON CORP (US)
                                                                                                                                                                                                                             /mol_type="unassigned DNA"
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                                                                                                                                                                                                                                                                                          /note="N = BIOTINYLATED C"
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/wol_type="unassigned DNA"
                                                                                                                                                                                                            /organism="unidentified"
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unidentified
unclassified.
1 (bases 1 to 20)
Atkinson, B.M. and Kealey, J.T.
PURIFIED TELOMERASE
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AX019595
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BD225828/c
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CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
0S Artificial Sequence
N JP 200250959-A/49
PD 02-APR-2002
PF 29-JAN-1999 JP 200529424
PR 29-JAN-1999 GB 9801902.9
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                                             PN 35 2001509681-A/7
PN 24-JUL-2001
PP 24-JUL-2001
PP 04-APR-1012-2001
PP 04-APR-1012-2001
PP 1 ALAIN PURSEROY, RONALD A PRUZAN, JAMES T KEALEY PC
C12N15/54, C12N9/12, C07K16/40, C12Q1/68, C07K14/47 CC Strandedness:
Single,
CC 7 Topology: Linear;
CC 7 Mod_base OTHER
CC / note= '0 ligonucleotide 14ab'
FF Key
Location/Qualifiers
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PC A61P35/00, C12N1/15, C12N1/19, C12N1/21, C12NS/10, C12P21/02 PC A61P35/00, C12N1/15, C12N1/19, C12N1/21, C12N3/10, C12N3/10, C12N3/10, A61K35/16, A61K31:522), C12N15/00, A61K37/02, C12N5/00 CC Description of Artificial Sequence:Oligonucleotide FH Key Location/Qualifiers
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100.0%; Pred. No. 1.3e+02;
tive 0; Mismatches 0;
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/mol_type="genomic DNA"
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Purified telomerase
Patent: JP 2001509681-A 7 24-JUL-2001;
GERON CORP
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Promoter region of mouse and BD225846
BD225846.1 GI:33035616
JP 2002509699-A/49.
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/organism="Zea mays"
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synthetic construct
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Best Local Similarity 90.9°
Matches 20, Conservative
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Best Local Similarity
Matches 19; Conserv
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PR 29-JAN-1998 GB 9801902.9
PI WILLIAM NICOL KEITH
PC
C12N15/09,A61K31/7105,A61K31/711,A61K35/76,A61K38/00,A61K45/00, PC
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human telomerase RNA component genes.
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PC A61P35/00, C12N1/15, C12N1/19, C12N1/21, C12N5/10, C12P21/02 .
C1201/68/(C12N9/12),
PC (A61K35/76, A61K31:522), C12N15/00, A61K37/02, C12N5/00 CC Description of Artificial Sequence: Primer FR Key

FT source / Organism='Artificial Sequence'.
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
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/organism='Artificial Sequence'
Location/Qualifiers
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/note="Oligonuclectide"
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JP 2002509699-A/31.
Synthetic construct
synthetic construct
other sequences; artificial sequences.
1 (bases 1 to 20)
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other sequences; artificial sequences.
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Sequence 49 from Patent WO9938964.
AX019595
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29-JAN-1999 JP 2000529424
29-JAN-1998 GB 9801902.9
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Best Local Similarity 90.94
Matches 20; Conservative
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US09436060A-16.rge.sl

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4.18;
95.08;
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Best Local Similarity 95.0
Matches 19; Conservative
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AX019580/c
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BD196339
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A61K48/00,
PC
A61P35/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02 PC
C12Q1/68/C12N9/12,
PC
C461K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00 CC
Description of Artificial Sequence: Primer
Location/Qualifiers
                                                                                                                                                                                                                                                                                                               Promoter region of mouse and human telomerase RNA component genes Patent: JP 2002509699-A 34 02-ARR-2002;
CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
OS Artificial Sequence
PN JP 2002509699-A/34
PP 02-ARR-2002
PP 29-JAN-1999 JP 2000529424
PR 29-JAN-1999 GB 9801902.9
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Promoter region of mouse and human telomerase RNA component genes.
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Patent: WO 9938964-A 31 05-AUG-1999;
KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4.1%; Score 18.4; DB 1; Length 20; 95.0%; Pred. No. 1.4e+02; tive 0; Mismatches 1; Indels
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Score 18.4; DB 1; Length 20;
Pred. No. 1.4e+02;
0; Mismatches 1; Indels
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/organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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synthetic construct
souther sequences; artificial sequences.
(bases 1 to 20)
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Sequence 31 from Patent WO9938964.
AX019577
                                                           102 TTCTCGCTGACTTTCAGCGG 121
20 TTCTCGCTGACTTCCAGCGG 1
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BD225831.1 GI:33035601
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    4.1%;
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synthetic construct
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Best Local Similarity 95.0%
   Query Match
Best Local Similarity 95.0
Matches 19; Conservative
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AX019577/c
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AUTHORS
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synthetic construct

synthetic construct

other sequences; artificial sequences.

the construct

other sequences; artificial sequences.

I (Dases 1 to 18)

Kilian, A. and Bowtell, D.

Kilian, A. and Bowtell, D.

Artificial Sequence

Synthetic and Systems LLC, PETER MACCALLUM CANCER INSTITUTE

Artificial Sequence

No Artificial Sequence

No JP 2002514928-A/73

PP 21-MAY-2002

PP 11-MAY-2002

PP 01-ULL-1997 US 60/053129,04-AUG-1997 US 60/054642 PR

21-ULL-1997 US 60/053329,04-AUG-1997 US 60/054642 PR

PC 11-ULL-1997 US 60/053329,04-AUG-1997 US 60/054642 PR

PC C12N15/54,C12N9/12,A61K38/45,C07K16/40,C12Q1/68,C12Q1/48, PC

C12N15/11,

PC A61K31/70

CC Description of Artificial Sequence:Synthesized Amplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Vertebrate telomerase genes and proteins and uses thereof. BD196339 BD196339.1 GI:33006109 UP 2002514928-A/73.
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
                                                                                                                Length 20;
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                                                                                                             Query Match
4.1%; Score 18.4; DB 1;
Best Local Similarity 95.0%; Pred. No. 1.4e+02;
Matches 19; Conservative 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 18.4; DB 1;
Pred. No. 1.4e+02;
0; Mismatches 1;
/organism="synthetic construct"
|mol_type="unassigned DNA"
|db_xref="taxon:32630"
|note="primer"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="synthetic construct"
|mol type="unassigned DNA"
|bxref="taxon:32630"
|note="primer"
                                                                                                                                                                                                                                                                                                                                                                             DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         other sequences; artificial sequences.
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Sequence 34 from Patent WO9938964.
AX019580
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C12N15/09,A61K31/7105,A61K31/711,A61K35/76,A61K38/00,A61K45/00,PC
                                                                                                                                               Promoter region of mouse and human telomerase RNA component genes batent: JP 2002509699-A 47 02-APR-2002;
CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
OS Artificial Sequence
PN JP 2002509699-A/47
PD 02-APR-2002
PP 29-JAN-1999 JP 2006529424
PR 29-JAN-1999 GB 9801902.9
PI WILLIAM NICOL KEITH
 Promoter region of mouse and human telomerase RNA component genes
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PC A61P35/00,C12N1/15,C12N1/19,C12N1/21,C12NS/10,C12P21/02 PC
,C12Q1/68//C12N9/12,
PC (A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00 CC
Description of Artificial Sequence:Oligonucleotide FH Key
Location/Qualifiers
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
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86.4%; Pred. No. 1.9e+02;
tive 0; Mismatches 3;
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3.8%; Score 17.2; DB 1;
Best Local Similarity 86.4%; Pred. No. 1.9e+02;
Matches 19; Conservative 0; Mismatches 3;

    .22
    ^coganism="synthetic construct"
/mol_type="genomic DNA"
    /db_xref="taxon:32630"

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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/noTe="Oligonucleotide"
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                                                                                                    other sequences; artificial sequences.
1 (bases 1 to 22)
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Sequence 47 from Patent WO9938964.
AX019593
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                                   BD225844.1 GI:33035614 JP 2002509699-A/47.
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                                                                    synthetic construct
synthetic construct
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synthetic construct
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Best Local Similarity 86.4'
Matches 19; Conservative
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DEFINITION
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SOURCE
ORGANISM
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AX019593
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SOURCE
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linkages are replaced by N-(2-aminoethyl)glycine units linked
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JP 2001517929-A/9
09-OCT-2001
09-ARR-1997 JP 1997536487
09-ARR-1996 US 08/630019
JERRY W SHAY,WOODRING E WRIGHT,MIECZYSLAW A PIATYSZEK,DAVID
                                                                                                                                                                                                                                                                                                                                                                                                                        BD071043.1 GI:22616646
JP 2001517929-A/9.
unidentified
unidentified
unclassified.
1 (bases 1 to 18)
Shay, J.W., Wright, W.E., Piatyszek, M.A., Corey, D. and Norton, J.C.
Modulation of mammalian telomerase by peptide nucleic acids
Patent: JP 2001517929-A 9 09-OCT-2001;
   Primer Design
based on EST Sequence GenBank Accession Number AA281296 FH
Location/Qualifiers
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                                                                    'organism='Artificial Sequence'
                                                                                                                                                                                   4.0%; Score 18; DB 1; Length 18;
100.0%; Pred. No. 1.4e+02;
tive 0; Mismatches 0; Indels
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Pred. No. 1.4e+02;
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                                                                                                    1. .18
/organism="synthetic construct"
/mol_type="genomic DNA"
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100.0%; Pred. No. ..

    .18
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Strandedness: Single;
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                                                                                  Location/Qualifiers
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                                                                                                                                                                      Query Match
Best Local Similarity 100.0
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JAMES C NORTON
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Best Local Similarity 100.0
Matches 18; Conservative
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DEFINITION
                                                                                                                                                                                                                                                                                                                                          RESULT 202
BD071043/c
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AUTHORS
TITLE
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VERSION
KEYWORDS
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ò g RESULT 207

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C1201/02, C1201/68, G01N33/15, G01N33/48, G01N33/50//C07K14/47, PC C07K16/40,
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                       PAT 18-JUN-2001
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C12N15/09, A61K31/70, A61K38/55, A61K39/395, A61K39/395, A61K48/00,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 pc (C12N1/21, C12R1:19), (C12N9/12, C12P21/08, (C12N1/19, PC (C12N1/21, C12R1:19), (C12N1/21, C12R1:84), PC (C12N9/12, C12R1:91), C12N15/00, A61R37/64, C12N5/00 CC Strandedness: Single; CC Topology: Linear; FR Key Location/Qualifiers FT Source 1.18
                                                                                                                                                                                                                                    unclassified.

1 (bases 1 to 18)

2 Thomas, R.S., Occhimu, R., Toru, N., Karen, B.C., Greg, B.M.,

Calvin, B.H. and William, H.A.

Human telomerase catalytic subunit promoter

L patent: JP 1999253177-A 201 21-SEP-1999;

URCORP, UNIVERSITY TECHNOLOGY CORP

OS Unidentified

PN JP 1999253177-A/201

PD 21-SEP-1999

PF 15-OCT-1996 US 08/724,643,18-APR-1997 US 08/841.419, Pl

25-APR-1997 US 08/846.017,06-MAY-1997 US 08/851.843, PR

14-AUG-1997 US 08/854.050,14-AUG-1997 US 08/915.503 PI TH

R SECHI, UCCHIMU RINGER, TORU MAXAMURA, KAREN B CHAPMAN, PI GREM

MORIN, PILLIAM, PILLIAM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 08/844.419,
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                           linear
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White the control of the c
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/organism='Unidentified'
Location/Qualifiers
                       E36993 18 bp DNA l
Human telomerase catalytic subunit promoter.
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Pred. No. 1.7e+02;
0; Mismatches 1;
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Sequence 543 from patent US 6610839.
AR390670
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                                                                                                                E36993.1 GI:13022956
JP 1999253177-A/201.
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Best Local Similarity 94.4%;
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Kim, W., W., W., W., W., W. Kealey, J.T., Pruzan, R. and Weinrich, S.L.
Method for detecting and inhibiting RNA component of telomerase
patent: JP 2001507229-A 14 05-JUN-2001;
GERON CORP
PN JP 2001507229-A/14
PD 05-JUN-2001
PP 19-DEC-1997 JP 1998529003
PP 20-DEC-1996 US 08/770564, Z0-DEC-1996 US 08/770565 PI
NAM WOO KIM, FRED WU, JAMES T KEALEY, RONALD PRUZAN, SCOTT L PI
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Methods for detecting the RNA component of telomerase
Patent: US 5846723.A 14 08-DEC-1998;
Location/Qualifiers
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PC C12N5/10,
C C12N9/12,C12Q1/68,C12Q1/68,C12N15/00,C12N5/00 CC
Strandedness: Single;
CC Topology: Linear;
CT Topology: Linear;
FH Key Location/Qualifiers.
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100.0%; Pred. No. 1.5e+02;
tive 0; Mismatches 0; Indels
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larity 100.0%; Pred. No. 1.5e+02;
Conservative 0; Mismatches 0;
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/organism="unknown"
/mol_type="unassigned DNA"
                                                                                            AR063838 17 bp 1 Sequence 14 from patent US 5846723.

    .17
    /organism="unidentified"
    /mol_type="genomic DNA"
    /db_xref="taxon:32644"

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JP 2001507229-A/14.
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THOMAS GREG B

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UN Vertebrate telomerase genes and proteins and uses thereof.

No Vertebrate telomerase genes and proteins and uses thereof.

No BD196340.

BD196340.

S JP 2002514928-A/74.

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SKILIan, A and Bowtell, D.

Vertebrate telomerase genes and proteins and uses thereof

RAILAN, A and Bowtell, D.

Vertebrate telomerase genes and proteins and uses thereof

Patent: JP 2002514928-A 74 21-MAY-2002;

CAMBIA BLOSYSTRAS LLC, PETER MACCALLUM CANCER INSTITUTE

OS Artificial Sequence

PN JP 2002514928-A/74

PD 21-MAY-2002

PR 01-JUL-1997 US 60/051410,21-JUL-1997 US 60/053018 PR

21-JUL-1997 US 60/053329,04-AUG-1997 US 60/054642 PR
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GREG B
        PAT 31-JAN-2002
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AGINSB/00, A61F35/00, A61F43/00,
COTKS/10,
COTKS/110,
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                                                                                                           DEVISION 101081042-A/201.

unidentified
unclassified.
1 (bases 1 to 18)
Sechi, T.R., Lingner, J., Nakamura, T., Chapman, K.B., Mori, G.B.,
Human telomerase catalytic subunit
Patent: JP 2001081042-A 201 27-MAR-2001,
GERON. CORP, UNIVERSITY TECHNOLOGY CORP
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CC Topology: Linear;
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PN JP 2001081042-A/201
PD 27-MAR-2001
PP 27-MAR-2001
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PP 02-CT-1996 US 08/724643, 18-APR-1997 US 08/84441
25-APR-1997 US 08/724643, 18-APR-1997 US 08/851843 PR 09-MAY-1997 US 08/854050, 14-AUG-1997 US 08/91312 PR 14-AUG-1997 US 08/912951, 14-AUG-1997 US 08/915503 PI R SECHI, JOACHIM LINGNER, TORU NAKAMURA, KAREN B CHAPMAN, PI MORIN.
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Human telomerase catalytic subunit.
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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                                                                                       BD011244.1 GI:18639617
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Best Local Similarity 94.4
Matches 17; Conservative
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Cech.T.R., Lingmer.J., Nakamura,T., Chapman,K.B., Morin,G.B.,
Harley,C.B. and Andrews,W.H.
Cells immortalized with telomerase reverse transcriptase for use in
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Harley,C.B. and Andrews,W.H.
Human telomerase catalytic subunit
Patent: EP 1313094-A 543 06-AUG-2003;
Geron Corporation (US); University Technology Corporation (US)
Location/Qualifiers
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3.6%; Score 16.4; DB 1; Length 18;
Best Local Similarity 94.4%; Pred. No. 1.7e+02;
Matches 17; Conservative 0; Mismatches 1; Indels
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3.6%; Score 16.4; DB 1; Length 18;
Best Local Similarity 94.4%; Pred. No. 1.7e+02;
Matches 17; Conservative 0; Mismatches 1; Indels
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Patent: US 6617110-A 543 09-SEP-2003;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                  Sequence 543 from patent US 6617110.

    .18
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    /db_xref="taxon:32644"

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/organism="unknown"
/mol_type="genomic DNA"
                                                                                                                                                 149 CACCGTTCATTCTAGAGC 166
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bp DNA linear PAT 17-JUL-2003
human telomerase RNA component genes.
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DN 4P 2002509699-A/17
PD 02-APR-2002
PF 29-JAN-1999 JP 2000529424
PR 29-JAN-1998 GB 9801902.9
PI WILLIAM NICOL KEITH
PC C12N15/09.A61K31/7105,A61K35/76,A61K38/00,A61K45/00, PC
                                                                                                                                                                                   linkages are replaced by N-(2-aminoethyl)glycine units linked
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CANCER RESPARCH CAMPAIGN TECHNOLOGY LTD
OS Artificial Sequence
PN JP 2002509699-A/17
PP 29-JAN-1999 JP 2000529424
PR 29-JAN-1999 GB 9801902.9
                                                                                                                   Strandedness: Single;
Topology: Linear;
/desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC
              09-APR-1997 JP 1997536487
09-APR-1996 US 08/630019
JERRY W SHAY, WOODRING E WRIGHT, MIECZYSLAW A PIATYSZEK, DAVID
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C A61P35/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02 PC
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PC (A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00
Description of Artificial Sequence: Primer
FH Key Location/Qualifiers
FT source 1. 20
Organism='Artificial Sequence'.
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0
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CC cc cc nucleotide bases via glycine amino N through a methylenecarbonyl linker'
FF Key location/Qualifiers
FT source longanism='Unidentified'.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism='Artificial Sequence' Location/Qualifiers
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/db_xref="taxon:32630"
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                                                                  COREY,
JAMES C NORTON
C07K14/00,A61K38/16,C12Q1/68
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/organism="unidentified"
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and
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synthetic construct
synthetic construct
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Best Local Similarity 100.0
Matches 16; Conservative
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Gorman, C.M. and Groskreutz, D.J.
Non-endocrine animal host cells capable of expressing variant proinsulin and processing the same to form active, mature insulin and methods of culturing such cells
Patent: US 6348327-A 13 19-FBB-2002;
Location/Qualifiers
                   ANDRZEJ KILIAN, DAVID BOWTELL
CI2N15/54, CI2N9/12, A61K38/45, C07K16/40, C12Q1/68, C12Q1/48, PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      l (bases 1 to 16)
Shay, "W, Wright, W.E., Piatyszek, M.A., Corey, D. and Norton, J.C. Modulation of mammalian telomerase by peptide nucleic acids Patent: JP 2001517929-A 26 09-OCT-2001; GERON CORP OS Unidentified
                                                                                    Description of Artificial Sequence:Synthesized Amplification
                                                                                                      Primer Design
based on EST Sequence GenBank Accession Number AA281296 FH
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Modulation of mammalian telomerase by peptide nucleic acids
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/mol_type="unassigned DNA"
                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                0; Mismatches
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   60/058287
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JP 2001517929-A/26.
unidentified
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Best Local Similarity 85.7
Matches 18; Conservative
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Best Local Similarity 85.7
Matches 18; Conservative
09-SEP-1997 US
PI ANDEZEJ KII
PC CL2N15/11,
PC A61K31/70
CC Description
CC Dased on EG
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FT source
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Bougueleret, L.
Nucleic acid encoding a retinoblastoma binding protein (RBP-7) and polymorphic markers associated with said nucleic acid Patent: US 6399373-A 76 04-JUN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PAT 22-JAN-2001
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LAX.-J. and Li, S.-H.
Hybridoma call line and monoclonal antibody for huntingtin protein
Patent: US 6355481-A 11 12-MAR-2002;
                                                              Gape
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85.0%; Pred. No. 2.3e+02;
ative 0; Mismatches 3; Indels
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                             Length 20;
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                           Query Match
3.4%; Score 15.4; DB 1;
Best Local Similarity 94.1%; Pred. No. 2.2e+02;
Matches 16; Conservative 0; Mismatches 1;
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Sequence 76 from patent US 6399373.
AR211866
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AR199735
AR199735.1 GI:20249809
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Sequence 11 from Patent WO0078813.
AX060355
/db_xref="taxon:9606"
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Best Local Similarity 85.0
Matches 17; Conservative
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AX060355
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AR199735
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20 bp DNA linear PAT 17-JUL-2003
Nucleic acid encoding retinoblastoma-binding protein (RBP-7) and
polymorphic marker relating to the nucleic acid.
BD221936
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JP 2002519027-A/75.
Homo sapiens (human)
Homo sapiens
Eukaryota: Metazos; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 20)
Bouguelere; L.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C12N15/09, C12N15/09, A01K67/027, C07K14/47, C07K16/18, C12N5/10,
                                                                                                                                                                                                                                                                                               Promoter regions of the mouse and human telomerase rna component
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           60/111909 PI
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PC C12N15/09.C12N15/09,A01K67/027,C07K14/47,C07K16/18,C12N5/PC C1201/68,
PC G01N33/53,G01N33/566,C12N15/00,C12N5/00,C12N15/00 CC Upstream amplification primer for SEQ 34, SEQ 55, SEQ 35, SEQ
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                                                                                                                                                                                                                                                                                                                              Pacent: WO 9938964-A 17 05-AUG-1999;
KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3.5%; Score 16; DB 1; Length 20;
100.0%; Pred. No. 2e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                linear
                                                                                                                                                                                                                                                                                                                                                                               1. 20
/organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/note="primer"
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                                                                                                                AX019563 20 bp
Sequence 17 from Patent WO9938964.
AX019563
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/organism="Homo sapiens"
/mol_type="genomic DNA"
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02-JUJ-2002
30-JUN-1999 JP 2000557360
30-JUN-1998 US 60/091315
IE BOUGUELERET
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                 CTCGGCTCACACATGC 451
                                      CTCGGCTCACACATGC 16
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Best Local Similarity 100.
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BD071036 15 bp DNA linear PAT 27-AUG-2002 Modilation of mammalian telomerase by peptide nucleic acids.
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unidentified
unidentified
unclassified.
1 (bases 1 to 15)
Shay,J.W., Wright,W.E., Piatyszek,M.A., Corey,D. and Norton,J.C.
Patent: JP 2001517929-A 2 09-OCT-2001;
GERON CORP
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JP 2001517929-A/2
09-0CT-2010
09-0CT-2010
09-APR-1996 US 08/630019
JERRY W SHAY,WOODRING E WRIGHT,MIECZYSLAW A PIATYSZEK,DAVID COREY,
COREY,
COTX14/00,AGIK38/16,C12Q1/68
Strandedness: Single;
            PR 20-DEC-1996 US 08/770564,20-DEC-1996 US 08/770565 PI NAM WOO KIM, FRED WU, JAMES T KEALEY, RONALD PRUZAN, SCOTT L PI
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                      CC linkages are replaced by N-(2-aminoethyl)glycin

CC cc nucleotide bases via glycine amino N through a

methylenecarbonyl linker'

FR Key Location/Qualifiers

1.15

/organism='Unidentified'.
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/organism='Unidentified'
Location/Qualifiers
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/organism="unidentified"
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/db_xref="taxon:32644"
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'organism="unidentified"
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/db_xref="taxon:32644"
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JP 2001517929-A/2.
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                                                                                                                                                               Huntington disease cellular model: stably transfected pc12 cells expressing mutant huntingtin Patent: WO 0079813-A 11 28-DEC-2000; Emory University (US) Location/Qualifiers
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Kim.N.W., Wu.F., Kealey, J.T., Pruzan, R. and Weinrich, S.L.

Method for detecting and inhibiting RNA component of telomerase Patent: JP 2001507229-A 10 05-JUN-2001;

PN JP 2001507229-A/10

PD 05-JUN-2001

PP 19-DP 10-DP 
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Unclassified.
Unclassified.
Kim,N.Woo., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L. Wethods for detecting the RNA component of telomerase Patent: US 5846723-A 10 08-DEC-1998;
Location/Qualifiers
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3.3%; Score 15; DB 1; Lv
Best Local Similarity 100.0%; Pred. No. 1.8e+02;
Matches 15; Conservative 0; Mismatches 0;

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    /organism="synthetic construct"

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                                                                                                   other sequences, artificial sequences
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/db_xref="taxon:32630"
/note="PCR primer"
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Sequence 10 from patent US 5846723,
AR063834
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/organism="unknown"
/mol_type="unassigned DNA"
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Modulation of mammalian telomerase by peptide nucleic acide
Modulation of mammalian telomerase by peptide nucleic acide

Datent: JP 2001517929-A 44 09-OCT-2001;
GGRON CORP
OS Unidentified

N JP 2001517929-A/44

PD 09-OCT-2001

PR 09-APR-1997 JP 1997536487

PR 09-APR-1996 US 08/630019

PI JERRY W SHAY, WOODRING E WRIGHT, MIECZYSLAW A PIATYSZEK, DAVID

PI JAMES C NORTON

PC COTXL4/00, A61K38/16, C12Q1/68

CC Strandedness: Single;

CC TOPOLOGY: Linear;

CC AGesc = 'phosphorothioate (PS) nucleic acid'

FH Key

Location/Qualifiers

FT SOURCE

/organism='Unidentified'.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PAT 27-AUG-2002
                                                                 Topology: Linear; //desc = 'poptide nucleic acid (PNA), where (deoxy(ribose- CC phosphate phosphate linkages are replaced by N-(2-aminoethyl)glycine units linked
                                                                                                                                                                                                                                                                                                                                                                                    Gape
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BD071078
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                                                                                                             CC linkages are replaced by N-(2-aminoethyl)glycin.
CC cc cc nucleotide bases via glycine amino N through a methylenecarbonyl linker'
FF Rey location/Qualifiers
FT source location/Oualified'.
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                                                                                                                                                                                                                    /organism='Unidentified'
Location/Qualifiers
              JAMES C NORTON
C07K14/00,A61K38/16,C12Q1/68
Strandedness: Single;
                                                                                                                                                                                                                                                          1. .15
/organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"

    .15
    /organism="unidentified"
    /mol_type="genomic DNA"
    /db_xref="taxon:32644"

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                                                                                                                                                                                                                                                                                                                                                                                                                                         1 ACCCTAACTGAGAAG 15
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                                                                                                                                                                                                                                                                                                                                                                                                                     49 ACCCTAACTGAGAAG 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CTAACCCTAACTGAG 1
                                                                                                                                                                                                                                                                                                                                                                                  15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 unclassified.
1 (bases 1 to 15)
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unidentified
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Best Local Similarity
Matches 15; Conserv
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Query Match
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DEFINITION
ACCESSION
VERSION
SCHWCES
SOURCE
ORGANISM
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BD071078/c
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TITLE
JOURNAL
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JP 2001517929-A/27.  
unidentified
                                                                 BD071039 15-bp DNA linear PAT 27-AUG-2002 Modulation of mammalian telomerase by peptide nucleic acids.
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JAMES C NORTON
COTK14/00, A61K38/16, C12Q1/68
Strandedness: Single;
Topology: Linear;
/desc = 'Peride nucleic acid (FNA), where (deoxy(ribose- CC
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JP 2001517929-A/27
09-OCT-2001
09-PR-1997 JP 1997536487
09-ARR-1996 US 08/630019
JERRY W SHAY,WOODRING E WRIGHT, MIECZYSLAW A PIATYSZEK, DAVID
                                                                                                                                                                                    unclassified.

1 (bases 1 to 15)
Shay, J.W., Wright, W.E., Piatyszek, M.A., Corey, D. and Norton, J.C. Modulation of mammalian telomerase by peptide nucleic acids Patent: JP 2001517929-A 5 09-OCT-2001;
                                                                                                                                                                                                                                                                                                                                                                  09-APR-1996 US 08/630019
JERRY W SHAY,WOODRING E WRIGHT,MIECZYSLAW A PIATYSZEK,DAVID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      unidentified
unclassified.
1 (bases 1 to 15)
Shay,J.W., Wright,W.E., Piatyszek,M.A., Corey,D. and Norton,J.C.
Modulation of mammalian telomerase by peptide nucleic acids
Patent: JP 2001517929-A 27 09-OCT-2001;
GERON CORP
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CC CC CC CC CC nucleotide bases via glycine amino N through a methylenecarbonyl linker' Location/Qualifiers 1.15 gource 1.15 /organism='Unidentified'.
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Location/Qualifiers
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100.0%; Pred. No. 1.8e+02;
tive 0; Mismatches 0;

    .15
    /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:3264"

                                                                                                                                                                                                                                                                                           Unidentified
JP 2001517929-A/S
09-OCT-2010
09-ARR-1997 JP 1997536487
09-APR-1996 US 08/630019
                                                                                                              DDC/1039.1 GI:22616642
JP 2001517929-A/5.
unidentified
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Best Local Similarity
Matches 15; Conserva
                                                                                                                                                                         unidentified
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PP
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PP
                              RESULT 224
BD071039/c
LOCUS
DEFINITION
ACCESSION
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SOURCE
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TITLE
JOURNAL
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Gaps

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SYN 26-MAR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1. .19
//organism="synthetic construct"
//mol type="other DNA"
//db_xref="taxon:32630"
//db_xref="forward primer for Japanese flounder microsatellite sequence Poli104MHFS"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PAT 17-JUL-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fuji,K., Kobayashi,K., Mizuta,A., Hasegawa,O., Tabata,K., Sakamoto,T. and Okamoto,N.
A genetic linkage map of the Japanease Flounder, (Paralichthys olivaceus) Unpublished
2 (bases 1 to 19)
Mizuta,A., Tabata,K., Kobayashi,K., Fuji,K., Sakamoto,T. and Okamoto,N.
                                                                                                                                                                                                                                                                                                            Synthetic construct DNA, forward primer for Japanese flounder microsatellite sequence Poli104HFS-M.
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                                                                                                                      Length 19;
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                                                                                                                                                                                                                                                                                              linear
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88.9%; Pred. No. 2.3e+02;
tive 0; Mismatches 2;
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synthetic construct
other sequences; artificial sequences.
1 (bases 1 to 18)
Kusu, M., Qiu,G. and Hunfrees,R.
CANCER CELL VACCINE
acid differentiation
Patent: US 5688643-A 22 18-NOV-1997;
Location/Qualifiers
                                                                   /organism="unknown"
/mol_type="unassigned DNA"
                                                                                                                                                                                                                                                                                           19 bp
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                                                                                                                                                                             348 GTTCAGGCCTTTCAGGCC 365
                                                                                                                                                                                                  89 CCCCCCCCTGTTTTCTC 106
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JP 2002531582-A/23.
                                                                                                                                                                                                                                                                                                                                                                                         synthetic construct
synthetic construct
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BD274798
                                                                                                               Query Match 3.3
Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Direct Submission
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Best Local Similarity
Matches 16; Conserv
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                                                     source
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AB175197
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ORGANISM
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KEYWORDS
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VERSION
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AUTHORS
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                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unknows:
Unclassified.
1 (Dases I to 18)
Schafer,M.P. and Reid T.M.
Rapid and sensitive method for detecting histoplasma capsulatum
Rapid Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 (Dases I to 19)
Oka,T., Matsunaga,H. and Yamane,A.
Method of nucleic acid-differentiation and assay kit for nucleic
                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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0
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                                             linear
                                                                                                                                                                                                                                                                                                                                                                       Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
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3.3%; Score 15; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 2.3e+02;
Matches 15; Conservative 0; Mismatches 0;
                                          AX613673 20 bp DNA
Sequence 4698 from Patent WO02072882.
                                                                                                                                                                                                                  Coronary chip
Patent: WO 02072882-A 4698 19-SEP-2002;
OGHAM GmbH (DE)
                                                                                                                                                                                                                                                                                    1. .20
/organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 4 from patent US 6469156.
AR241369
AR241369.1 GI:27287033
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Sequence 22 from patent US 5688643.
174343 GI:3010484
                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                   Cullen, P. and Seedorf, U.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   410 CTGAGCTGTGGGACGTGC 427
                                                                                       AX613673.1 GI:28409102
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                                                                                                                                                                                                                                                                                                                                                                                                                                269 GGGCTTCTCCGGAGG 283
                                                                                                                                                                                                                                                                                                                                                                                                                                                               3 édéctroroceases 17
                                                                                                                       Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 3.39
Best Local Similarity 88.99
Matches 16; Conservative
                                                                                                                                        Homo sapiens
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Unclassified.
                                                                           AX613673
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                                                    DEFINITION
ACCESSION
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SOURCE
ORGANISM
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           RESULT 227
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AUTHORS
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AR241369/c
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VERSION
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TITLE
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                         AX613673
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Voss,J. and Timm,J.
Methods and compositions for the screening of cell cycle modulators
Patent: WO 0120034-A 17 22-MAR-2001;
BASF ARTENGESELSCHAFT (DE)
Location/Qualifiers
   INNOGENETICS N.V. (BE) ; Enterprise Ireland (trading as Bioresearch
Ireland) (IE)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mus musculus
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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1 (bases 1 to 17)

Andersson,L., Georges,M., Spincemaille,G. and Nezer,C.D.A.
Selection of animal based on character imprinted by parent patent: JP 2002535963-A 130 29-OCT-2002;
UNIVERSITY OF LIEGE,MELICA HB, SEGHERS GENTEC NV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PC C12N15/09,A01K67/027,C12N5/06,C12Q1/68,C12N15/00,C12NS/00 CC /note='Polymorphism Tyrosine Hydroxylase gene' FH Key
                                                                                                                                                                                     Gарв
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JP 2002535963-A/130
29-CCT-2002
16-DEC-1999 JP 2000588390
16-DEC-1998 EP 98204291.3
LEIF ANDERSSON,MICHEL GEORGES,GEERT SPINCEMAILLE, PI
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                                                                         /organism="Aspergillus versicolor"
/mol_type="unassigned DNA"
/db_xref="taxon:46472"
                                                                                                                                                Query Match
3.1%; Score 14; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 2.4e+02;
Matches 14; Conservative 0; Mismatches 0;
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100.0%; Pred. No. 2.4e+02;
tive 0; Mismatches 0;
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/mol_type="unassigned DNA"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                           Sequence 17 from Patent W00120034. AX099957
                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                         Mus musculus (house mouse)
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Selection of animal based
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BD251610.1 GI:33061380
JP 2002535963-A/130.
Sus scrofa (pig)
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Best Local Similarity 100.0%
Matches 14; Conservative
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Aspergillus versicolor
Eukaryota; Fungi; Ascomycota; Pezizomycotina; Eurotiomycetes;
Eurotiales; Trichocomaceae; mitosporic Trichocomaceae; Aspergillus.
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Patent: JP 2002531582-A 23 24-SEP-2002;
ANTIGEN EXPRESS INC
S. Artificial Sequence
PN JP 2002531582-A/23
PD 24-SEP-2002
PF 24-NOV-1999 JP 2000586901
PF 04-NOV-1999 JP 09/205995
PF 04-DEC-1998 US 09/205995
PI minzhen kusu,gang qiu,robert hunfrees
CC Description of Artificial Sequence: antisense oligonucleotide
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Xu,M., Qiu,G. and Humphreys,R.
MHC class II antigen presenting cells containing oligonucleotides
which inhibit II protein expression
Patent: US 636855-A 2 09-APR-2002;
Location/Qualifiers
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Weide,M.
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                                                                                                                                                               corresponding
to a specific region of the mouse Ii gene.
Key
Location/Qualifiers
                                                                                                                                                                                                                                        1. .18
/organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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Patent: WO 0073499-A 21 07-DEC-2000;
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/organism="unknown"
/mol_type="unassigned DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AR205264 18 bp 18 Sequence 24 from patent US 6368855.
AR205264
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Best Local Similarity 93.84
Matches 15; Conservative
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Unclassified.
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PAT 16-SEP-2000
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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Butheria, Primates, Catarrhini, Hominidae, Homo.
                                                                                                                                                                                    Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Cetartiodactyla, Suina, Suidae, Sus.
                                                                                                                                                                                                                              Georges, M., Spincemaille, G. and Andersson, L. Selecting animals for parentally imprinted traits Patent: WO 035143-A 130 22-UN-2000, SEGHENSGENTEC N V (BE); GEORGES MICHEL (BE); UNIV LIEGE (BE) SPINCEMAILE GEERT (BE); MELICA HB (SE); ANDERSSON LEIF (SE) Location/Qualifiers
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Method and reagent for the inhibition of grid
Patent: WO 0162911-A 129 30-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); GLAXO GROUP LIMITED
Location/Qualifiers
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/organism="Sus scrofa"
/mol_type="unassigned DNA"
/morfe="taxon:9823"
/noFe="Polymorphism Tyrosine Hydroxylase gene"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
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AX272560.1 GI:16545297
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Sequence 330 from Patent WO0162911.
AX272761
AX272761.1 GI:16545498
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/mol_type="unassigned RNA"
/db xref="taxon:9606"
                                                                                  Sequence 130 from Patent WO0036143.
                                                                    17 bp
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                                                                                                     AX028311
AX028311.1 GI:10189199
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3.1%;
Best Local Similarity 88.2%;
Matches 15; Conservative
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                                                                                                                                                     Sus scrofa (pig)
                                                                                                                                                                      Sus scrofa
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AX272761/c
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AX028311
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Zwick,M.G., Edington,B.E., McSwiggen,J.A., Merlo,P.Ann.Owens.,
Zwick,M.G., Skokut,T.A., Young,S.A., Folkerts,O. and Merlo,D.J.
Nucleic acid encoding delta-9 desaturase
Patent: US 6350934-A 826 26-FEB-2002;
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                                                                                                                                                     ch 3.1%; Score 13.8; DB 1; Length 17; I Similarity 88.2%; Pred. No. 2.4e+02; 15; Conservative 0; Mismatches 2; Indels
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                                 /organism='Sus scrofa (pig)'. Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                           linear
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Seguence 826 from patent US 6350934.
AR196361
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PNA-DNA-PNA chimeric macromolecules
Patent: US 5700922-A 2 23-DEC-1997;
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="unknown"
/mol_type="unassigned DNA"
                                                                                                                                                                                                                                                                                                                                      17 bp
Sequence 2 from patent US 5700922.
186370
                                                                       1. .17
/organism="Sus scrofa"
/mol_type="genomic DNA"
/db_xref="taxon:9823"
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Location/Qualifiers
FT source
FT
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Best Local Similarity 88.2
Matches 15; Conservative
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Unclassified.
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Matches 15; Conserv
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186370
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Length 17;

PAT 31-MAR-2003

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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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3.1%; Score 13.8; DB 1;
Best Local Similarity 88.2%; Pred. No. 2.4e+02;
Matches 15; Conservative 0; Mismatches 2;
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3.1%; Score 13.8; DB 1;
Best Local Similarity 88.2%; Pred. No. 2.4e+02;
Matches 15; Conservative 0; Mismatches 2;
                  1..17
/organism="synthetic construct"
/mol_type="unassigned DNA"
/db xref="taxon:32630"
/note="PNA analogue"
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Molecular Engines Laboratories (FR)
Location/Qualifiers
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Sequence 5862 from Patent WO03025176.
AX728175
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Sequence 814 from Patent EP1281758.
AX688082
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/mol_type="unassigned DNA"
/db_xref="taxon:10090"
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/db_xref="taxon:9606"
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Location/Qualifiers
  Location/Qualifiers
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Mus musculus
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                                                        Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Randi, A. M.
Randi, A. M.
Method and reagent for the inhibition of erg
Patent: WO OBB124-A 1949 22-NOV-2001,
RIBOZYME PHRMACEUTICALS, INC. (US); GLAXO GROUP LIMITED (GB)
Location/Qualifiers
                                                                                                              Jarvis, T., von Carlowitz, I., Mcswiggen, J.A., Hamblin, P.A. and Ellis, J.H.
Method and reagent for the inhibition of grid
Patent: WO 162911-A 330 30-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); GLAXO GROUP LIMITED (GB)
Location/Qualifiers
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88.2%; Pred. No. 2.4e+02;
iive 0; Mismatches 2; Indels
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3.1%; Score 13.8; DB 1; Length 17;
Best Local Similarity 88.2%; Pred. No. 2.4e+02;
Matches 15; Conservative 0; Mismatches 2; Indels
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synthetic construct
other sequences; artificial sequences.
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Sequence 1949 from Patent WO0188124.
AX423613
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Pna-dna-pna chimeric macromolecules
Patent: EP 1201676-A 2 02-MAY-2002;
PSISS PHARMACEUTICALS, INC. (US)
                                                                                                                                                                                                                                                   /organism="Homo sapiens"
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/db_xref="taxon:9606"

    .17
    /organism="Homo sapiens"
    /mol_type="unassigned RNA"
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Sequence 2 from Patent EP1201676.
AX429297.1 GI:21540603
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                    sapiens (human)
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Best Local Similarity 88.2<sup>5</sup>
....hes 15, Conservative
                                        Homo sapiens
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AX429297
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AX423613/c
                                      ORGANISM
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Length 17;

PAT 08-MAY-2003

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PAT 07-SEP-2000
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Guida, M. and Hall, J.
Genetic typing of the human cytochrome P450 2A6 gene and related
Genetic typing of the human cytochrome P450 2A6 gene and related
materials and methods
Patent: US 6492115-A 11 10-DEC-2002;
Location/Qualifiers
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Patent: WO 9938964-A 40 05-AUG-1999;
KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
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                       Length 38;
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                     Query Match
3.1%; Score 13.8; DB 1;
Best Local Similarity 72.0%; Pred. No. 3.2e+02;
Matches 18; Conservative 0; Mismatches 7;
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Pred. No. 3.2e+02;
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/organism="synthetic construct"
/mol_type="unassigned DNA"
/db xref="teaxon:32630"
/noTe="Oligonucleotide"
                                                                                                                                                                                                                                                                                                      DNA
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Sequence 11 from patent US 6492115.
AR264866.1 GI:29693235
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Sequence 40 from Patent WO9938964.
AX019586
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/organism="unknown"
/wol_type="genomic DNA"
                                                                                                                       222 TCGCCTGCCCAGCCCCCGAACCCCG 246
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Unclassified.
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AR264867
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CI2N15/09,A61K31/7105,A61K31/711,A61K35/76,A61K38/00,A61K45/00, PC
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Promoter region of mouse and human telomerase RNA component genes.
BD225837
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CANCER RESEARCH CAMPAGN TECHNOLOGY LTD
OS Attificial Sequence
PN JP 2002509699-A/40
PD 02-ARP-2002
PR 29-JAN-1999 JP 2000529424
PR 29-JAN-1998 GB 9801902.9
PI WILLIAM NICOL KEITH
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PC A61P35/00.C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02
C12Q1/68//C12N9/12,
CC (A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00 CC
Description of Artificial Sequence:Oligonucleotide FH Key
Location/Qualifiers
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                          Indels
88.2%; Pred. No. 2.4e+02; ive 0; Mismatches 2;

    .38
        /organism="synthetic construct"
/mol type="genomic DNA"
/db_xref="taxon:32630"

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Sequence 571 from patent US 6350934.
AR196106.
AR196106.1 GI:20245543
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/organism="unknown"
/mol_type="unassigned DNA"
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                                                                          368 AGGAAGAGGAACGGAGC 384
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JP 2002509699-A/40.
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                                                                                                                          17 AGGAAAAGGAACGGATC 1
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Best Local Similarity 88.2
Matches 15; Conservative
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BD225837/c
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Method and reagent for the inhibition of grid
Parent: WO 0162911-A 857 30-AUG-2001;
RIBOZYME PHARMACETICALS, INC. (US); GLAXO GROUP LIMITED (GB)
LOCATION/Qualifiers
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Patent: WO 0188124-A 1481 22-NOV-2001;
RIBOZYME PHARMACEUTICALS, INC. (US) ; GLAXO GROUP LIMITED (GB)
Location/Qualifiers
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Pred. No. 2.6e+02;
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.larity 93.3%; Pred. No. 2.6e+02;
Conservative 0; Mismatches 1;
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Sequence 1481 from Patent WO0188124.
AX423145.
AX423145.1 GI:21526527
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism="Homo sapiens"
/mol_type="unassigned RNA"
/db_xref="taxon:9606"
                   1. .17
/organism="Homo sapiens"
/mol_type="unassigned RNA"
/db_xref="taxon:9606"
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/mol_type="unassigned RNA"
/db xref="taxon:9606"
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Seguence 857 from Patent W00162911.
AX273288
 Location/Qualifiers
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Best Local Similarity 93.3%;
Matches 14; Conservative
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Homo sapiens
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AX423145/c
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                Unclassified.

1 (bases 1 to 17)

Guida, M. and Hall, J.

Genetic typing of the human cytochrome P450 2A6 gene and related materials and methods
Patent: US 6492115-A 12 10-DEC-2002;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unknown.
Unclassified.

Unclassified.

(bases 1 to 17)

Pavco, P., McSwiggen, J.A., Stinchcomb, D.T. and Escobedo, J.

Pavco, P., McSwiggen, for the treatment of diseases or conditions
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6566127-A 4559 20-MAY-2003;
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Method and reagent for the inhibition of grid
Patent: WO 0162911-A 429 30-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); GLAXO GROUP LIMITED (GB)
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Pred. No. 2.6e+02;
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US 6566127.
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AX272860
AX272860.1 GI:16545597
Sequence 12 from patent US 6492115.
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/wol_type="unassigned RNA"
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/organism="unknown"
/mol_type="genomic DNA"
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Sequence 4559 from patent
AR327157
AR327157.1 GI:33712965
                                                 AR264867.1 GI:29693236
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Best Local Similarity 93.3.
Best Local 4; Conservative
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Best Local Similarity 93.3
Matches 14; Conservative
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                                   AR264867
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Shay,J.W., Wright,W.E., Piatyszek,M.A., Corey,D. and Norton,J.C.
Modulation of mammalian telomerase by peptide nucleic acids
Patent: JP 2001517929-A 1 09-OCT-2001;
GERON CORP
                                                                                                                                                                                                                                                                                           Gaps
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Modulation of mammalian telomerase by peptide nucleic acids
                                                          Unclassified.

1 (bases 1 to 13)
Gryaznov,S., Pongracz,K. and Matray,T.
Oligonuclectide N3',Ewdarw.P5' thiophosphoramidates: their synthesis and administration to treat neoplasms
Patent: US 6608036-A 8 19-AUG-2003;
Location/Qualifiers
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/organism="unidentified"
/mol-type="unassigned RNA"
/db_xref="taxon:32644"
/note="This peptide is synthetically generated"
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2.9%; Score 13; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 2.2e+02;
Matches 13; Conservative 0; Mismatches 0;
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Reduction of halr growth
Patent: WO 03102077-A 3 09-JAN-2003;
THE GILLETTE COMPANY (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                             13 bp
Sequence 3 from Patent WO03002077.
                                                                                                                                                                                     1. .13
/organism="unknown"
/mol_type="genomic DNA"
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BD071035.1 GI:22616638
JP 2001517929-A/1.
unidentified
AR381412.1 GI:40089445
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JP 2001517929-A/1
09-0CT-2001
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Best Local Similarity 100.
Matches 13; Conservative
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BD071035/c
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                                                                                                                                                                                                                                                           Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Sequences involved in phenomena of tumour suppression, tumour reversion, apoptosis and/or resistance to viruses and the use thereof as medicaments and an analyse and an analyse and the use Patent: WO 3032117-A 1134 27-MAR-2003;
Molecular Engines Laboratories (FR)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unclassified.

1 (bases 1 to 13)

Gryaznov,S., Pongracz,K. and Matray,T.
Oligonucleotide N3'.fwdarw.P5' thiophosphoramidates: their synthesis and administration to treat neoplasms
Patent: US 6608036-A 2 19-AUG-2003;
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                                                                                                                                       linear
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2.9%; Score 13; DB 1; Length 13;
Best Local Similarity 100.0%; Pred. No. 2.2e+02;
Matches 13; Conservative 0; Mismatches 0; Indels
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Sequence 1134 from Patent W003025177.
AX735544
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                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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Sequence 2 from patent US 6608036.
AR381406
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Sequence 8 from patent US 6608036.
AR381412

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                   124 GGAAAAGCCTCGGCC 138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      97 TGTTTTTCTCGCTGA 111
                                       16 GGAAAAGCCTCCGCC 2
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BDU71079 13 bp DNA linear PAT 27-AUG-2002 Modulation of mammalian telomerase by peptide nucleic acids. BD071079
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Strandedness: Single;
Topology: Linear;
/desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC phosphate
linkages are replaced by N-(2-aminoethyl)glycine units linked
                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-APR-1996 US 08/630019
09-APR-1996 US 08/630019
JERRY W SHAY, WOODRING E WRIGHT, MIECZYSLAW A PIATYSZEK, DAVID
COREY,
                                                                                                                                                                                                                                                                                                                                      Shay, J.W., Wright, W.E., Piatyszek, M.A., Corey, D. and Norton, J.C. Modulation of mammalian telomerase by peptide nucleic acids Patent: JP 2001517929-A 12 09-OCT-2001;
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                                    Gape
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CC cc cclectide bases via glycine amino N through a methylenecarbonyl linker'
FH Key l. Ccation/Qualifiers
FT source l. 1. 13
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/organism='Unidentified'
Location/Qualifiers
Query Match
2.9%; Score 13; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 2.2e+02;
Matches 13; Conservative 0; Mismatches 0;
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2.9%; Score 13; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 2.2e+02;
Matches 13; Conservative 0; Mismatches 0;

    .13
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    /mol_type="genomic DNA"
    /db_xref="taxon:32644"

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OS Unidentified

PN JP 201517929-A/12

PD 09-CT-2001

PF 09-APR-1997 JP 19975:

PR 09-APR-1996 US 08,

PI JERRY W SHAY, WOODRING

PI JAMES C NORTON

PI JAMES C NORTON

CC Strandedness: Single, CC

CC TOPOLOGY: Linear;

CC TOPOLOGY: Linear;

CC TOPOLOGY: Linear;

CC TOPOLOGY: Linear;
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JP 2001517929-A/12.
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JP 2001517929-A/45.
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1 (bases 1 to 13)
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BD071079/c
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Modulation of mammalian telomerase by peptide nucleic acids.
BD071038
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                                                                                                                                                                     linkages are replaced by N-(2-aminoethyl)glycine units linked
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JAMES C NORTHALO, AGINSB, 16, C12Q1/68
Strandedness: Single;
Topology: Linear;
Jeptide nucleic acid (PNA), where (deoxy(ribose- CC phosphate
                                                                                                             Topology: Linear; //desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC phosphate
 09-APR-1997 JP 1997536487
09-APR-1996 US 08/630019
JERRY W SHAY, WOODRING E WRIGHT, MIECZYSLAW A PIATYSZEK, DAVID
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Shay,J.W., Wright,W.E., Piatyszek,M.A., Corey,D. and Norton,J.C. Modulation of mammalian telomerase by peptide nucleic acids. Pacent: JP 2001517929-A 4 09-OCT-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nucleotide bases via glycine amino N through a lenecarbonyl linker'
                                                                                                                                                                                                                                                                                                                                                                                        Query Match 2.9%; Score 13; DB 1; Length 13; Best Local Similarity 100.0%; Pred. No. 2.2e+02; Matches 13; Conservative 0; Mismatches 0; Indels
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Location/Qualifiers
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Location/Qualifiers
                                                                                                                                                                                                                                    Location/Qualifiers

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                                                  COREY,
JAMES C NORTON
CO7K14/00, A61K38/16,C12Q1/68
Strandedness: Single;

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JP 2001517929-A/4
09-OCT-2001
09-ARR-1997 JP 1997536487
09-ARR-1996 US 08/630019
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JP 2001517929-A/4.
unidentified
unidentified
unclassified.
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BD071038/c
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AUTHORS
TITLE
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BD071085 13 bp DNA linear PAT 27-AUG-2002 Modulation of mammalian telomerase by peptide nucleic acids. BD071085 BD071085 1 GI:22616688 UP 2001517929-A/51. unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Topology: Linear;
/desc = 'poptide nucleic acid (PNA), where (deoxy(ribose- CC
phosphate
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Shay,J.W., Wright,W.E., Piatyszek,M.A., Corey,D. and Norton,J.C. Modulation of mammalian telomerase by peptide nucleic acids Patent: JP 2001517929-A 51 09-OCT-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unidentified
JP 2001517929-A/51
09-0CT-2001
09-APR-1997 JP 1997536487
09-APR-1996 US 08/630019
JERRY W SHAY, WOODRING E WRIGHT, MIECZYSLAW A PIATYSZEK, DAVID
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unclassified.
1 (bases it to 13)
Shay,J.W., Wright,W.E., Piatyszek,M.A., Corey,D. and Norton,J.C.
Modulation of mammalian telomerase by peptide nucleic acids
Patent: JP 2001517929-A 54 09-OCT-2001;
GERON CORP
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CC cc nucleotide bases via glycine amino N through a

methylanecarbonyl linker'

FH Key location/Qualifiers

FT source loganism='Unidentified'.
                                 Indels
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Location/Qualifiers
         Pred. No. 2.2e+02;
; Mismatches 0;
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CO7K14/00,A61K38/16,C12Q1/68
Strandedness: Single;

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0
       Best Local Similarity 100.0%;
Matches 13; Conservative C
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Matches 13; Conservative
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unclassified.
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BD071085/c
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AUTHORS
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unclassified.

In Chases 1 to 13)
Shay, J.W., Wright, W.B., Piatyszek, M.A., Corey, D. and Norton, J.C.
Modulation of mammalian telomerase by peptide nucleic acids.

Patent: JP 2001517929-A 50 09-OCT-2001;
GERON CORP
OS Unidentified
PN JP 2001517929-A/50
PD 09-OCT-2001
PP 09-APR-1997 JP 1997536487
PR 09-APR-1996 US 08/630019
PI JERRY W SHAY, WOODRING E WRIGHT, MIECZYSLAW A PIATYSZEK, DAVID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Modulation of mammalian telomerase by peptide nucleic acids.
BD071084.1 GI:22616687
UP SO0151739-A/50.
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linkages are replaced by N-(2-aminoethyl)glycine units linked
                   Unidentified
JP 2001517929-A/45
09-OCT-2001
09-OCT-2001
09-APR-1997 JP 1997536487
09-APR-1996 US 08/630019
JERRY W SHAY, WOODRING E WRIGHT, MIECZYSLAW A PIATYSZEK, DAVID
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JAMES C NORTON
COTK14/00.AGIK18/16,C12Q1/68
COTK14/00.AGIK18/16,C12Q1/68
COTCAGOGES: Single;
C Topology: Linear;
C Topology: Linear;
C Topology: Linear;
Location/Qualifiers
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                        /organism='Unidentified'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 2.9%; Score 13; DB 1; Le Best Local Similarity 100.0%; Pred. No. 2.2e+02; Matches 13; Conservative 0; Mismatches 0;

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Strandedness: Single;

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JAMES C NORTON
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GERON OS UN 
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BD071084/c
LOCUS
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ACCESSION
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SOURCE
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TITLE
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PAT 17-AUG-2003
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(C12NS/00,C12R1:91)
Regulation of repressor genes using nucleic acid molecules FH
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PF 11-APR-2000 JP 2000611654
PR 12-APR-1999 US 60/129390
PI LAWRENCE BLATT MICHAEL ZWICK, PAWELA PAVCO, JAMES MCSWIGGEN PC C12N15/09, A61K38/00, A61K48/00, A61P43/00, A61P43/00, C12N5/10, PC C12P21/02,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (C12P21/02, C12R1:91), (C12P21/02, C12R1:91), C12N15/00, C12N5/00,
                                                                                                                                                                                                                                                                                                                                        Tobases 1 to 16)
Paul, P.S., Meng, X.-J., Halbur, P., Morozov, I. and Lum, M.A.
Polymucleic acids isolated from a porcine reproductive and
respiratory syndrome virus (PRRSV) and proteins encoded by the
polymucleic acids
Patent: US 6592873-A 34 15-JUL-2003;
                                                                                Gape
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BD253907
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Blatt, L., Zwick, M., Pavco, P. and Mcswiggen, J.
Blatt, L., Zwick, M., Pavco, P. and Mcswiggen, J.
Bregulation of represeor genes using nucleic acid molecules
Patent: JP 2002541795-A 1700 10-DEC-2002;
RIBOZYME PHARMACEUTICALS INC
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                                                  Length 13;
                                                                                                                                                                                                                       linear
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                                                Query Match
2.9%; Score 13; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 2.2e+02;
Matches 13; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                       DNA
                                                                                                                                                                                                               AR353144
Sequence 34 from patent US 6592873.
AR353144
    /mol_type="genomic DNA"
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    16
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JP 2002541795-A/1700
10-DEC-2002
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JP 2002541795-A/1700.
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                                                                                                            55 ACTGAGAAGGGCG 67
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unclassified.
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AUTHORS
TITLE
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linkages are replaced by N-(2-aminoethyl)glycine units linked
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                                                                                                                                                         Topology: Linear; /desc = 'peptide nucleic acid (PNA), where (deoxy(ribose- CC
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Unidentified
JP 2001517929-A/54
09-0CT-2001
09-0CT-201
09-APR-1995 US 08/630019
JERRY W SHAY.WOODRING E WRIGHT, MIECZYSLAW A PIATYSZEK, DAVID COREY.
COREY
COREY
COREY
CORTON
COTKIA/00, AGIK38/16, C12Q1/68
Strandedness: Single;
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JP 2001537929-A/55
0-OCT-2001
09-APR-1997 JP 1997536487
09-APR-1996 US 08/630019
JERRY W SHAY,WOODRING E WRIGHT,MIECZYSLAW A PIATYSZEK,DAVID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 (bases 1 to 13)
Shay J.W., Wright, W.E., Piatyszek, M.A., Corey, D. and Norton, J.C. Modulation of mammalian telomerase by peptide nucleic acids Patent: JP 2001517929-A 55 09-OCT-2001;
GERON CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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Modulation of mammalian telomerase by peptide nucleic acids
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                                                                                                                                                                                        2.9%; Score 13; DB 1; Length 13; 100.0%; Pred. No. 2.2e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                              /organism='Unidentified'
Location/Qualifiers
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/organism='Unidentified'
Location/Qualifiers

    .13
    /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"

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JAMES C NORTON
CO7K14/00,A61K38/16,C12Q1/68
Strandedness: Single;
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BD071089.1 GI:22616692
JP 2001517929-A/55.
unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        53 TAACTGAGAAGGG 65
                                                                                                                                                                                                                                                                                                                                                                                                                                           13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          unidentified unclassified.
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unclassified.

unclassified.

l (bases 1 to 17)

S Blatt.i., Zwick,M., Pavco,P. and Mcswiggen,J.

Regulation of repressor genes using nucleic acid molecules

RIBOZYME PHARMACEUTICALS INC

OS BURATYOLE

PN JP 2002541795-A/6149

PD 10-DEC-2002

PP 11-APR-2000

PP 11-APR-2000

PP 12-AR-1999 US 60/129390

PI 1AMYBNCE BLATT,MICHAEL ZWICK,PAMELA PAVCO,JAMES MCSWIGGEN PC

CIZNIS/09,A61K38/00,A61K48/00,A61P43/00,A61P43/00,CIZNS/10, PC
                  (bases 1 to 17)
Blasczyk,R.D.
A method for determining the Histocompatibility locus antigen class
                                                                                                                                                                                                                                                                                                                                                                                                                        BUZSB356 17 bp DNA linear PAT 17-JUL-2003
Regulation of repressor genes using nucleic acid molecules.
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2.8%; Score 12.8, DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 2.8e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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ilarity 87.5%; Pred. No. 2.8e+02;
Conservative 0; Mismatches 2:
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/organism="unidentified"
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                                                                                  Patent: EP 0887423-A 3 30-DEC-1998;
BIOTEST AG (DE)
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/organism="unidentified"
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BD258356.1 GI:33068126
JP 2002541795-A/6149.
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BD273168
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BD258356
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C12Q1/68, C12M1/00, C12N15/09, C12N15/09, C12N15/09, G01N33/53, PC
                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic construct
synthetic construct
construct
coher sequences; artificial sequences.
1 (bases 1 to 16)
2 Fan, J.B., Hirschhorn, J.N., Huang, X., Kaplan, P., Lander, E.S.,
Lockhart, D.J., Ryder, T. and Sklar, P.
Universal arrays
WHITEHEAD INSTITUTE FOR BIOMEDICAL RESEARCH, AFFYMETRIX INC
OS Artificial Sequence
PN JP 2002539849-A/444
PD 26-NOV-2002
PF 77-MAR-2000 JP 2000608794
PR 26-MAR-1999 US 60/140359 PI
JIAN BING FAN, JOEL N HIRSCHHORN, XIAOHUA
HUANG, PAUL KAPLAN, ERIC
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2.8%; Score 12.8; DB 1; Length 16;
Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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Pred. No. 2.7e+02;
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PI DAVID J LOCKHART, THOMAS RYDER, PAMELA SKLAR
PC C1201/68,C12M1/00,C12N15/09,C12N15/09,C12N15/00
G01N33/566,
PC G01N37/00,C12N15/00,C12N15/00,C12N15/00
CC Primer Location/Qualifiers
FH Key Location/Qualifiers
FT Source /organism='Artificial Seque
                          /organism='Eukaryote'.
Location/Qualifiers
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/organism="synthetic construct"
/mol_type="genomic DNA"
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Sequence 3 from Patent BP0887423.
A82083.
A82083.1 GI:6731948
                                                                          /organism="unidentified"
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           1. .17
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JP 2002539849-A/444.
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A82083
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TITLE

8 셤 COMMENT

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Expression vectors comprising multiple shear stress responsive elements (SSRE) and methods of use for treating disorders related to vasculogenesis and/or angiogenesis in a shear stress environment Patent: US 6440726-A 1 27-AUG-2002;

Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 2.8%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 2.8e+02; Matches 14; Conservative 0; Mismatches 2; Indels
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87.5%; Pred. No. 2.8e+02;
ative 0; Mismatches 2;
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/mol_type="unassigned DNA"
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BIOTEST AG (DE)
Location/Qualifiers
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                                                                                                                                         AR224417 17 bp
Sequence 1 from patent US 6440726.
AR224417
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Sequence 3 from Patent EP0892069.
AX001154.1 GI:7241348
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                   GCTGACTTTCAGCGGG
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1 (bases 1 to 17)
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1 (bases 1 to 17)
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Matches 14; Conservative
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AR224417
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OS Artificial Sequence

NO 4002533113-A/1

PD 08-0CT-2002

PF 23-DEC-1999 JP 2000591168

PR 24-DEC-1998 US 60/113863,24-DEC-1998 US 09/220510 PI

NITZAN RESNICK

PC C12N15/09,A61K31/713,A61K35/76,A61K48/00,A61P3/06,A61P3/10, PC
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PC C12N1/15,
PC C12N1/15,C12N1/10,G01N33/15,G01N33/50,C12N15/00,C12N5/ PC
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Expression vectors comprising multiple shear stress responsive elements (SSRE) and a gene of interest and methods of use thereof.
                                                                                                                                                                          Expression vectors comprising multiple shear stress responsive elements (SSRE) and a gene of interest and methods of use thereof Patent: JP 2002533113-A 1 08-OCT-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                    A61P11/00, A61P13/10, A61P17/02, A61P19/02, A61P35/00, A61P43/00, C12N1/15,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Description of Artificial sequence: A PDGF-A Shear Stress CC
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Unknown.
Unclassified.
I (bases 1 to 17)
Zwick,M.G., Edington,B.E., McSwiggen,J.A., Merlo,P.Ann.Owens.,Guo,L., Skokut,T.A., Young,S.A., Folkerts,O. and Merlo,D.J.
Nucleic acid encoding delta-9 desaturase
Patent: US 6350934-A 828 26-FEB-2002;
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism='Artificial Sequence'
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/organism="synthetic construct"
/mol type="genomic DNA"
/db_xref="taxon:32630"
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                                                                                                                       other sequences, artificial sequences.
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AR196363.1 GI:20245800
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/mol_type="unassigned DNA"
                                                    BD273168.1 GI:33082936
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AR196363
   DEFINITION
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TITLE
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Fattaey, A.R., Jarvis, T., Mcswiggen, J., Booher, R.N. and Holman, P.S. Method and reagent for the inhibition of checkpoint kinase-1 (chk
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Method and reagent for the modulation and diagnosis of cd20 and nogo gene expression
Patent: WO 0159103-A 1810 16-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); Blatt, Lawrence (US);
McSwiggen, James (US); Chowrira, Bharat M. (US)
Location/Qualifiers
          nogo gene expression
Patent: WO 0159103-A 59 16-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); Blatt, Lawrence (US)
McSwiggen, James (US); Chowrira, Bharat M. (US)
Location/Qualifiers
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Patent: WO 0157206-A 7 09-AUG-2001;
RIBOZYWE PHARMACEUTICALS, INC. (US) ; Fattaey, Ali R. (US)
Location/Qualifiers
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/organism="synthetic construct"
/mol_type="unassigned RNA"
/db_xref="taxon:32630"
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Seguence 7 from Patent W00157206.
AX226635
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KEYWORDS
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                                                                                              Blatt, L., Mcswiggen, J. and Chowrira, B.M.
Method and reagent for the modulation and diagnosis of cd20 and nogo gene expression
Patent: WO 01531013-A 27 16-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); Blatt, Lawrence (US);
McSwiggen, James (US); Chowrira, Bharat M. (US)
Location/Qualifiers
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Patent: WO 0159103-A S8 16-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); Blatt, Lawrence (US)
McSwiggen, James (US); Chowrira, Bharat M. (US)
Location/Qualifiers
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/mol type="unassigned RNA"
/mol xref="taxon:32630"
/nofe="Nucleic Acid"
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/mol_type="unassigned RNA"
/db_xref="taxon:32630"
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                                                              other sequences; artificial sequences.
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synthetic construct
other sequences; artificial sequences.
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Sequence 58 from Patent W00159103.
AX214616
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AX214617/c
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ORGANISM
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VERSION
KEYWORDS
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TITLE
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TITLE
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                                                                                                                                                                                                                                                                                        Kmiec, E.B., Gamper, H.B., Rice, M.C. and Kim, J.
Targeted chromosomal genomic alterations in plants using modified single stranded oligonucleotides
Patent: WO 0195512-A 722 06-DEC-2001;
UNIVERSITY OF DELAWARE (US)
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Method and reagent for the inhibition of erg
Patent: WO 0188124-A 818 22-NOV-2001;
RIBOZYME PHARMACEUTICALS, INC. (US) ; GLAXO GROUP LIMITED (GB)
Location/Qualifiers
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Sequence 838 from Patent WO0188124.
AX422502
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Sequence 839 from Patent WO0188124.
AX422503
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Sequence 752 from Patent W00192512.
AX324614
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/mol_type="unassigned DNA"
/db_xref="taxon:4577"

    .17
/organism="Homo sapiens"
/mol_type="unassigned RNA"
/db_xref="taxon:9606"

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CTGAGCTGAGGGCCGT 16
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Best Local Similarity 87.5'
Matches 14; Conservative
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AX422503/c
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Spermatophyta, Magnoliophyta, Liliopsida, Poales, Poaceae, PACCAD
clade, Panicoideae, Andropogoneae, Zea.
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Targeted chromosomal genomic alterations in plants using modified single stranded obligonucleotides
Patent: WO 0195512-A 751 06-DEC-2001;
UNIVERSITY OF DELAWARE (US)
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Method and reagent for the inhibition of grid
Patent: WO 0162911-A 331 30-AUG-2001;
RIBOZYME PHARMACHICALS, INC. (US) ; GLAXO GROUP LIMITED (GB)
Location/Qualifiers
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87.5%; Pred. No. 2.8e+02;
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2.8%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 2.8e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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    .17
    /organism="Homo sapiens"
    /mol_type="unassigned RNA"
    /db_xref="taxon:9606"

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AX324613
AX324613.1 GI:18095366
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/db_xref="taxon:32630"
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/organism="Zea mays"
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AX324613
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PAT 26-NOV-2002

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Homo sapiens
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Human udp-galnac:polypeptide n-acetylgalatosaminyltransferase 10

Patent: EP 1243660-A 739 25-SEP-2002;

Aeomica, Inc. (US)

Location/Qualifiers
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Zhang,J., Gu,Y. and Nguyen,C.T.
Human udp-galnac:polypeptide n-acetylgalatosaminyltransferase 10
Patent: EP 1243660-A 738 25-SEP-2002;
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2.8%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 2.8e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                                                        ch 2.8%; Score 12.8; DB 1; Length 17; ll Similarity 87.5%; Pred. No. 2.8e+02; 14; Conservative 0; Mismatches 2; Indels
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2.8%; Score 12.8; DB 1;
Best Local Similarity 87.5%; Pred. No. 2.8e+02;
Matches 14; Conservative 0; Mismatches 2;
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/mol_type="unassigned DNA"
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Sequence 739 from Patent EP1243660.
AX545226

    .17
/organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"

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Sequence 738 from Patent EP1243660.
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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Homo sapiens
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Inc. (US)
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Best Local Similarity
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 Aeomica,
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AX545226/c
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AX545225/c
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                             Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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                                                                                                                                Jarvis,T., von Carlowitz,I., Mcswiggen,J.A., Mclaughlin,F.G. a Randi,A.M.
Method and reagent for the inhibition of erg
Patent: WO 0188124-A 839 22-NOV-2001;
RIBOZYME PHARMACEUTICALS, INC. (US) ; GLAXO GROUP LIMITED (GB)
Location/Qualifiers
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Human posh-like protein 1
Patent: EP 1239051-A 716 11-SEP-2002;
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Patent: BP 1239051-A 715 11-SEP-2002;
Aeomica, Inc. (US)
Location/Qualifiers
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Sequence 716 from Patent EP1239051.
AX531207.
AX531207.1 GI:25254207
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned RNA"
/db_xref="taxon:9606"
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       AX422503.1 GI:21525885
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PAT 26-NOV-2002

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Kit and method for determining HLA type
Patent: WO 019257-2A 1071 06-DEC-2001;
NISSHINDO INDUSTRIES INC. SYSTEM RESEARCH INC, HIDETOSHI INOKO, TAEKO
KAGIYA, TATSUO ICHIHARA, YOSHIYUKI MATSUMURA, SHOGO MORIYA, MICHIO
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Patent: JP 2002509699-A 44 02-APR-2002;
CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
OS Artificial Sequence
PN JP 2002509699-A/44
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                                 synthetic construct
synthetic construct
other sequences; artificial sequences.
1 (bases 1 to 17)
Incko, H., Kagiya, T., Ichihara, T., Matsumura, Y., Moriya, S. and
Nishida, M.
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C12Q1/68,C12M1/00,C12N15/09,G01N33/53
Description of Artificial Sequence:capture
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/organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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other sequences; artificial sequences.
1 (bases 1 to 25)
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WO 0192572-A/1071
06-DEC-2001
01-JUN-2001 WO 2001JP0004662
01-JUN-2000 JP 00P 164798
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29-JAN-1998 GB 9801902.9
WILLIAM NICOL KEITH
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BD104967.1 GI:22650541
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PF 01-JUN
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MATSUMURA,
PI SHOGO
PC C1201/,
CC Descri
FH KEY
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and mdz12
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Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7
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2.8%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 2.8e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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Kit and method for determining HLA type.
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Aeomica, Inc. (US)
Location/Qualifiers
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Aeomica, Inc. (US)
Location/Qualifiers
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Sequence 815 from Patent EP1281758.
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Sequence 813 from Patent EP1281758.
                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/db_xref="taxon:9606"
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/organism="Homo sapiens"
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 343 GCGAGGTTCAGGCCTT 358
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                                 16 GCGCGGATCAGGCCTT
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BD104967/c
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BD023698 30 bp DNA linear PAT 27-AUG-2002 Method for detecting and inhibiting RNA component of telomerase.
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OS Artificial Sequence
NO JP 2002505699-A/42
PD 02-APR-2002
PP 29-JAN-1999 GB 9801902.9
PP 29-JAN-1999 GB 9801902.9
PI WILLIAM NICOL KEITH
C12NIS/09, A61X31/7105, A61X31/711, A61X35/76, A61X38/00, A61X45/00, PC
A61X48/00.
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Promoter region of mouse and human telomerase RNA component genes.
BD225839
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                                                                                                                                                                                                                        unidentified unclassified.

unclassified.

1 (bases 1 to 30)

Kim,N.W., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.

Method for detecting and inhibiting RNA component of telomerase Patent: JP 2001507229-A 2 05-JUN-2001;

GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                      PN JP 2001507229-A/2
PD 05-UIN-2001
PF 19-DEC-1997 JP 1998529003
PR 20-DEC-1996 US 08/770564,20-DEC-1996 US 08/770565 PI
NAM WOO KIM, FRED WU, JAMES T KEALEY, RONALD PRUZAN, SCOTT L PI
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PC C12N15/09, A61K9/08, A61K31/7105, A61K45/00, A61K48/0
PC C12N5/10,
PC Topology: Linear;
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Pred. No. 3.6e+02;
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1 (bases 1 to 38)
Keith, W.N.
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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JP 2001507229-A/2.
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Best Local Similarity 70.8%;
Matches 17; Conservative
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Kim,N.Woo., Wu,F., Kealey,J.T., Pruzan,R. and Weinrich,S.L.
Methods for detecting the RNA component of telomerase
Patent: US 5846723-A 2 08-DEC-1998;
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 12.8; DB 1; Length 25; Pred. No. 3.5e+02;
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                                                                                                                     Length 25;
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                                                                                                                     2.8%; Score 12.8; DB 1; 70.8%; Pred. No. 3.5e+02; tive 0; Mismatches 7;

    .25
    /organism="synthetic construct"
|mol_type="genomic DNA"
|db_xref="taxon:32630"

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/organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="teaxon:32630"
/noTe="Oligonucleotide"
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synthetic construct
other sequences; artificial sequences.
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Sequence 44 from Patent WO9938964.
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/organism="unknown"
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AX019590.1 GI:10043504
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Best Local Similarity 70.8%;
Matches 17; Conservative
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Unclassified.
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AR063826
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AX019590/c
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Roninson I.B. and Grossman, A.
Compositions, methods and kits for identifying naturally occurring
RNA sequences having affinity for RNA-binding proteins
Patent: US 6238867-A 15 29-MAY-2001;
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PR 29-JNN-1998 GB 9801902.9
PI WILLIAM NICOL KEITH
PC CI2N15/09, A61K31/7105, A61K31/711, A61K35/76, A61K38/00, A61K45/00, PC
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UP 2002509201
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CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
OS Artificial Sequence
PN JP 2002509699-A/83
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PC A61P35/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/02 PC,

C12Q1/68//C12N9/12,

PC (A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00 CC

Description of Artificial Sequence: Mutant construct FH Key

Location/Qualifiers
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                                                                                                                                   linear
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JP 20025096591-A/83.
synthetic construct
synthetic construct
other sequences; artificial sequences.
1 (bases 1 to 51)
                                                                                                                                AR154150 66 bp 1
Sequence 15 from patent US 6238867.
AR154150 GI:15122203
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/mol_type="unassigned DNA"
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29-JAN-1999 JP 2000529424
29-JAN-1998 GB 9801902.9
WILLIAM NICOL KEITH
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Unclassified.
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BD225880/c
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Roninson, I.B. and Grossman, A.
Compositions, methods and kits for identifying naturally occurring
RNA sequences having affinity for RNA-binding proteins
Patent: US 6238867-A 6 29-MAY-2001;
Location/Qualifiers
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
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                                                                                                                                                  Query Match 2.8%; Score 12.8; DB 1; Length 38; Best Local Similarity 70.8%; Pred. No. 3.3e+02; Matches 17; Conservative 0; Mismatches 7; Indels
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/noTe="Oligonucleotide"
                                                                 1. .38
/organism="synthetic construct"
/mol_type="genomic DNA"
/db_xrefe"taxon:32630"
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AR154151
AR154151.1 GI:15122204
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/mol_type="unassigned DNA"
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/organism="synthetic construct"
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/db_xref="taxon:32630"
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Matches 13; Conserv
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Promoter region of mouse and human telomerase RNA component genes.
BD225882
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CANCER RESEARCH CAMPAIGN TECHNOLOGY LTD
OS Artificial Sequence
PN JP 2002509699-A/85
PD 02-APR-2002
                                                                                                                                                                                                                                                                                    Keith, W.N. Promoter regions of the mouse and human telomerase rna component genes
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PC A61P35/00,C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21,C12Q1/68//C12N9/12,
PC (A61K35/76,A61K31:522),C12N15/00,A61K37/02,C12N5/00
Description of Artificial Sequence: Mutant construct FH
Location/Qualifiers
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   Pred. No. 2.7e+02;
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                                            263 GGCCCGGGGCTTCTCCGGAGGCACCCACTGCCACC 297
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60.0%; Pred. No. 2.7e+02;
tive 0; Mismatches 14;
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/organism="synthetic construct"
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/noTe="Mutant construct"
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Keith, W.N.
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                                                                                                                                                Sequence 83 from Patent W09938964.
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29-JAN-1998 GB 9801902.9
WILLIAM NICOL KEITH
 60.0%; Pred. No.
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Matches 21; Conservative
Best Local Similarity
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AX019629/c
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BD225882/c
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Sullivan, S., Dapper, K., Kisich, K., Stinchcomb, D.T. and McSwiggen, J.
TNF-.alpha. ribozyman, A. 58 22-SEP-1998;
Patent: Location/Qualifiers
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KEITH WILLIAM NICOL (GB); CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
Query Match
2.8%; Score 12.6; DB 1; Length 51;
Best Local Similarity 60.0%; Pred. No. 2.7e+02;
Matches 21; Conservative 0; Mismatches 14; Indels
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Best Local Similarity 60.0%; Pred. No. 2.7e+02;
Matches 21; Conservative 0; Mismatches 14; Indels
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/noTe="Mutant construct"
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AR041868
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Sequence 85 from Patent WO9938964.
AX019631
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/wol_type="unassigned DNA"
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/mol_type="genomic DNA"
  Location/Qualifiers
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Balzergue, S.
Direct Submission
Direct Submission
Submitted (11-SEP-2004) Balzergue S., UMRGV, INRA/CNRS, 2 rue
Gaston Cremieux, 91057 Evry cedex, FRANCE
PCR was performed on DNA from transforments of Arabidopsis thaliana
pCR was performed on DNA from transforment (s) resulting from
the PCR were directly sequence from the left or the right border
to determine the genomic sequence flanking the insertion. T-DNA
derived sequences were removed. Information to order the
corresponding mutant line and a link to a database providing a
graphical display of the insertion site are available at
http://dbagap.versailles.inra.fr/publiclines/. This sequence has
been generated in the framework of the French plant genomics
program 'Genoplante-info.infobiogen.fr).
                                         PAT 21-FEB-2003
                                                                                                                                                                                                                                                                                                                          Method and reagent for inhibiting the expression of disease related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ô
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T-DNA integration into the Arabidopsis genome depends on sequences of pre-insertion sites

EMBO Rep. 3 (12), 1152-1157 (2002)
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                                         linear
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RIBOZYME PHARMACEUTICALS, INC. (US)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     right border, T-DNA flanking sequence. Arabidopsis thaliana (thale cress) Arabidopsis thaliana
                               Sequence 4425 from Patent EP1260586. AX637286 AX637286.1 GI:28472900
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AX637286/c
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AJ840863
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                                                                              /clone="615B02"
/clone_lib="Arabidopsis thaliana T-DNA insertion lines"
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Method of diagnosing colon and gastric cancers
Patent: WO 2004021010-A 81 11-MAR-2004;
Oncotherapy Science, Inc. (JP); Japan as represented by the president of the university of Tokyo (JP)
Location/Qualifiers
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/noTe="Artificially synthesized S-oligonucleotide"
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Unclassified.
1 (bases 1 to 16)
Guida,M. and Kurth,J.
Genotyping the human phenol sulfotransferbase 2 gene STP2
Patent: US 6448003-A 38 10-SEP-2002;
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organism="Arabidopsis thaliana"
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/note="T-DNA flanking sequence
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Sequence 81 from Patent WO2004021010.
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                  /mol_type="genomic DNA"
/cultivar="Wassillewskija"
                                                            /db_xref="taxon:3702"
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Novel human gene relating to respiratory diseases, obesity, and inflammatory bowel disease

Litanmacory bowel disease

Location 0178894-A 316 25-OCT-2001;

Genome Therapeutics Corp. (US)

Location/Qualifiers

Location/Qualifiers

Location/Qualifiers

Location/Qualifiers

Amol type="unassigned DNA"

/db xref="taxon:32630"
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Query Match 2.7%; Score 12.4; DB 1; Length 16; Best Local Similarity 92.9%; Pred. No. 2.8e+02; Matches 13; Conservative 0; Mismatches 1; Indels
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GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen nucleic search, using sw model	st 24, 2005, 14:24:45;	US-09-436-060A-16 451 1 gggttgcggaggtgggcctagg	NTITY N OP 10.0	479 segs, 9939 residues f hits satisfying chosen parameters:	length: 0 length: 200000000	g: Minimum Match 0% Maximum Match 100% Listing first 488 summaries	rng.subdb:*	. is the number of results predicater than or equal to the score	SUMMARIES	% Query Match Length DB ID	17.5 79 1 12.0 62 1		10.2 51 1	8.8 8.8 47 1 8.1 47 1	8.0 38 1 6.9 38 1	6.9	6.9 31 1	6.9 6.9 31 11	6.7 30 1	6.7 30 1	6.7 30 1	6.7	6.7 30 1	6.7 30 1 6.7 30 1 6.7 30 1	6.7 30 1
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2.8 17 1 ADMS8919 Hepatitis B virus 2.8 17 1 ADI83671 HCV DNAzyme substr 2.8 17 1 ADI86393 HCV DNAzyme substr	8 17 1 ADI84179 HCV DNAZYME	8 17 1 AD186509 HCV DNAZYME 8 17 1 ADN44083 Mutant cell	17 1	8 1/1 ADQ80/40 FOLCINE 8 1/1 ADQ92762 Androger	8 25 1 AAZ07300	8 30 1 AAV41159 8 38 1 AAZ07298	8 62 1 AAA08205 Adenovirus nuc	8 62 1 AAH24816	8 66 L	8 15 1 ADG98448		OTHER PRINCIPLE	CINCIPAL			ADP27647 standard; DNA; 79 BP.			(first entry)		DNA used as a cancer prognostic marker SeqID 84.	: PCR amplicon; ds: prognostic marker; EGFR;	factor receptor; cancer; gene expression p	head and neck cancer; colon cancer; metastatic spread;	יד פעמפע.	ń	- A 1.			; 2003WO-US036777.		. 2002US-0427090P.	GENOMIC HEALTH INC.	VALL HEBRON UNIV HOSFITAL.	Cronin MT, Shak S, Baselga J;	2004-420643/39.			analysis	gene e.g., CD44v3.	400000000000000000000000000000000000000	Claim 55; SEQ ID NO 84; 113pp; English.	ion relates to a novel method concerning prognostic markers	with EGFR (epidermal growth factor receptor) positive cancer.	/, it refers to a gene expression profitting method that can rediction as to whether a natient is likely to respond well to	ith an EGFR inhibitor. The present invention describes the	analysis of the expression level of the RNA transcript of at	RRII7, LAMC2 or their products thereof. It further provides a CDNA	microarray containing named genes that represent prognostic transcripts	which are useful for determining whether a patient diagnosed with an EGFR -expressing head or neck cancer or colon cancer exhibits elevated or	decreased expression levels of these genes compared to normal. As such,
12.8	12.8	12.8	12.8	12.8	12.8	12.8	12.8	12.8	12.8	12.6						27647 sta		ADF21641;	26-AUG-2004		Human TERC I	C: human	epidermal growth		neoprantic o	o sapiens	WO2004046386-A1		03-JUN-2004	14-NOV-2003;		5-NOV-2002;		(VALL-) VALI	Baker JB, (, 2004-42			subjecting a quantitative	east one ge	, i	im 55; SE	s invent	ociated	ciricaliy wide a n	atment wi	ntitative	17, LAMC	roarray	ch are us pressing	reased ex
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Match 2'-O-methyl Telomerase inhibit Telomerase templat	Labelled nucleic a	Labelled nucleic a Human G/C-rich (Sp	Human B-raf target	Z',5'-Oligoadenyli Galanin receptor g	Human LCAT gene po	human CETF gene at PRRSV sequencing b		Human H-Ras DNAzym	8	tory olic	Serotonin receptor	Pyrimidine nucleot	Antiviral phosphor	Delta-9 desaturase	Human BRCA1 probe	Human leukocyte an		Human adenosine re	Hammerhead ribozym	Shear Stress Respo	Human Chki ribozym	Human NOGO Hammerh	Human NOGO Hammerh	Human NOGO Zinzyme	Human GRID NCH rib	Human pp-GaNTase 1	Human pp-GaNTase 1 Male-sterile plant	Male-sterile plant	Human ERG hammerhe	Human POSHLI scann	Human POSHL1 scann	Human HLA genotypı WNV minus strand I	WNV Zinzyme substr	WNV minus strand A WNV Hammerhead Rib	WNV Inozyme substr	WNV Minus Strang D Human MDZ3 scannin	Human MDZ3 scannin	Human H-Ras DNAzym	human hekz DNAzyme Human H-Ras DNAzym		Human H-Ras DNAzym	HCV DNAzyme substr	HCV minus strand D	HBV inozyme substr	SNX9 (Sorting Nexi		Human PTGDR substr	Human NOGO recepto	Human IKK-gamma su	Human GRID mRNA su Human C/EBPN DNA f	Human glioma endot
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invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  detecting a first subunit protein (1) of telomerase. R1 and R2 both bind to (1) and have formulae 5'-A-B-C-3' (R1) 5'-D-F-7' (R2) where: A and F c (1) and have formulae 5'-A-B-C-3' (R1) 5'-D-F-7' (R2) where: A and F c RNA segment of 10 to 100000 nucleotides (nt) that together are replicated by RNA replicase; B and E = RNA segments of 10 to 250 nt from the Y region of human telomerase and bind specifically to (1); C and D = RNA segments of 1 to 10000 nt which can be ligated together. Ligation of RNA segments of 1 to 10000 nt which can be ligated together. Ligation of to (1). Replication of R3 by RNA replicase indicates presence of (1). Also described are: (1) method for detecting (I) using R1 and R2; (2) kit for this process containing R1, R2, pigase and an amplification system; and (3) method for making R1 and R2 by transcription from appropriate DNA. R1 and R2 are used to detect and quantify telomerase, particularly for diagnosis of cancer and for detection of metastases. R1 and R2 provide an assay that does not require expensive equipment or highly trained personnel, and is suitable for automation. The present sequence represents an oligonucleotide used in the exemplification of the present
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these methods are also useful for prognosing or predicting the likelihood of cancer-attributable death or progression, including recurrence and metastatic spread of a neoplastic disease, as well as drug resistance. This polynuclectide sequence is a human PCR amplicon DNA sequence used a prognostic cancer marker, given in an exemplification of the invention.
                                                                                                                                                                           371 AAGAGGAACGGAGCGAGTCCCCGCGCGCGCGCGATTCCCTGAGCTGTGGGACGTGCACC 430
                                                                                                                                                                                                      1 AAGAGGAACGGAGTCCCCGCGCGCGCGCGCATTCCCTGAGCTGTGGGACGTGCACC 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pair of RNA molecules for detecting telomerase, useful for diagnosis of cancer or metastases, can be ligated when bound to telomerase subunit
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human, telomerase, hTR, telomeric repeat amplification protocol; TRAP,
identification, detection; quantification, cancer, metastasis, ss.
                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes a pair of RNA molecules (R1, R2) for
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0
                                                                                                                   Length 79;
                                                                                                                                              0; Indels
                                                                                      Sequence 79 BP; 16 A; 26 C; 27 G; 10 T; 0 U; 0 Other;
                                                                                                                 Score 79; DB 1; 1
Pred. No. 0.0059;
                                                                                                                                                                                                                                                                                                                                                                                                                           Adenovirus nucleotide sequence SEQ ID NO:20.
                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 2; Col 23; 32pp; English.
                                                                                                                                                                                                                                 431 CAGGACTCGGCTCACACAT 449
                                                                                                                                                                                                                                                   CAGGACTCGGCTCACACAT 79
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                                                                                                                   17.5%; S
100.0%;
                                                                                                                                                                                                                                                                                                                                       BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       97US-00873709
                                                                                                                                                                                                                                                                                                                                     AAA08205 standard; DNA; 62
                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                              79; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MPI; 2000-282223/24
                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mastadenovirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                               28-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14-MAR-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Grossman A;
                                                                                                                                                                                                                                                                                                                                                                   AAA08205;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein.
                                                                                                                                                                                                                                                                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New ribonucleic acids useful for identifying naturally occurring RNA sequences having affinity for RNA-binding protein having protein and RNA
                                                                                                                                                                                                                                                                                                               RNA-binding protein; RBP; RNA replicase; RNA identification; telomerase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The specification describes a first RNA (RNA1) and a second RNA (RNA2)
                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gapa
                                                                                       54
                                                                                                                   8
                                                                                                                                                                                                                                                                                   Human nucleic acid sequence derived from Y-1 domain of telomerase.
                                                                                                                    1 GGGTTGCGGAGGGTGGCCCTGGGAGGGGTGGTGGCCATTTTTGTCTAACCCTA
                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
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                               DB 1; Length 62;
0.7;
                                                           IndelB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
Sequence 62 BP; 16 A; 27 C; 10 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 62 BP; 16 A; 27 C; 10 G; 9 T; 0 U; 0 Other;
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                    12.0%; Scor.
100.0%; Pred. No. v. ...
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ilarity 100.0%; Pred. No. 0.7
Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (INVI-) INVITRO DIAGNOSTICS INC.
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                                                                                                                                                                                            AAH24816 standard; RNA; 62
                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Roninson IB, Grossman A;
                                                            54; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-366472/38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
es' 54; Conserv
                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                         US6238867-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-FEB-1998;
                                                                                                                                                                                                                                                       06-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                       29-MAY-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      components.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    telomerase
                                                                                                                                                                                                                         AAH24816;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                RESULT 3
AAH24816/c
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Matches'
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RNA-binding protein; RBP; RNA replicase; RNA identification; telomerase;

Homo sapiens.

US6238867-B1

(INVI-) INVITRO DIAGNOSTICS INC.

WPI; 2001-366472/38.

Roninson IB,

99US-00255464 98US-0075495P

22-FEB-1999; 23-FEB-1998;

29-MAY-2001

Human nucleic acid sequence derived from Y-1 domain of telomerase.

AAH24815 standard; RNA; 66 BP.

06-AUG-2001

AAH24815;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes a pair of RNA molecules (R1, R2) for detecting a first subunit protein (1) of telomerase. R1 and R2 both bind to (1) and have formulae 5'-A-B-C-3' (R1) 5'-D-E-F-3' (R2) where: A and F = RNA segment of 10 to 100000 mucleotides (nt) that together are replicated by RNA replicase; B and E = RNA segments of 10 to 250 nt from the Y region of human telomerase and bind specifically to (1); C and D = RNA segments of 1 to 10000 nt which can be ligated together. Ligation of C and D produces R3 of formula 5'-A-B-C-D-E-F-3' (R3) with E and B bound to (1). Replication of R3 by RNA replicase indicates presence of (1). Also described are: (1) method for detecting (1) using R1 and R2; (2) kit for this process containing R1, R2, ligase and an amplification system; and (3) method for making R1 and R2 by transcription from appropriate DNA. R1 and R2 are used to detect and quantify telomerase, particularly for diagnosis of cancer and for detection of metastases. R1 and R2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     provide an assay that does not require expensive equipment or highly trained personnel, and is suitable for automation. The present sequence represents an oligonucleotide used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pair of RNA molecules for detecting telomerase, useful for diagnosis of cancer or metastases, can be ligated when bound to telomerase subunit
                                                                                                                                                                                                                                                                                                                                                                                                                     Human; telomerase; hTR; telomeric repeat amplification protocol; TRAP; identification; detection; quantification; cancer; metastasis; ss.
                        54
                                                                    61 GGGTTGCGGAGGGTGGGCCTGGGAGGGGTGGTGGCCATTTTTTGTCTAACCCTA 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12.0%; Score 54; DB 1; Length 66; 100.0%; Pred. No. 0.77;
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                                                                                                                                                                                                                                                                                                                                                                    Adenovirus nucleotide sequence SEQ ID NO:19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 2; Col 23; 32pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (INVI-) INVITRO DIAGNOSTICS INC.
                                                                                                                                                                                                          AAA08204 standard; DNA; 66 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mastadenovirus
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                                                                                                                                                                                                                                                            AAA08204;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 protein.
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                                                                                                                                                                               RESULT
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The specification describes a first RNA (RNA1) and a second RNA (RNA2) for use in binding an RNA-binding protein (RBP) having protein and RNA components. RNA1 has the formula 5'-A-B-C-3', where A is section having (10-100,000 nucleotides and is can be received by an RNA replicase and with another DNA sequence, F, being replicated; B is section having 10-3,000 nucleotides which have affinity to one consensus sequence of the RBP and which can bind to the protein component; C is section having about 1-20 nucleotides and which can be ligated to D of the second RNA molecule. RNA2 has the formula 5'-D-E-F-3', where D is section having 1-C on nucleotides and which can be ligated to C; E is section having 1-C and which can bind to the protein component; F is section having 10-100,000 nucleotides which have affinity to second consensus sequence of the RBP and which can bind to the protein component; F is section having 10-100,000 nucleotides which and RNA2 are capable of forming a third RNA (RNA3) of formula 5'A-B-C-D-E-F-3'. The method is useful for the identification of confidential calfinity to anino and characterization of RNA sequences having specific affinity to anino
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
New ribonucleic acids useful for identifying naturally occurring RNA sequences having affinity for RNA-binding protein having protein and RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             acid consensus sequences of RBPs, and to RNAS. AAR24815-16 were used to produce a double-stranded RNA1, comprising the Y-1 domain of human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 GGGTTGCGGAGGCTGGGAGGGGTGGTGGCCATTTTTGTCTAACCCTA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GGTTGCGGAGGCTGGGCCTGGGAGGGTGGTGGCCATTTTTTGTCTAACCCTA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mutated hTR promoter fragment containing construct 29111 (mRCE).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12.0%; Score 54; DB 1; Length 66; 100.0%; Pred. No. 0.77;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 66 BP; 10 A; 11 C; 28 G; 17 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Pred. ....
                                                                                                                                                  Example 2; Col 26; 36pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAZ00334 standard; DNA; 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                     components.
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AAZ00334
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ð q RESULT 5 AAH24815

Best Local Similarity

54 59

1 GGGTTGCGGAGGGTGGGCCTGGGAGGGGGTGGTGGCCATTTTTTTGTCTAACCCTA

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WO9938964-A2
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  THE STATE OF THE S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to promoter regions from mouse and human telomerase

RNA (TR) component genes. The TR gene promoter can be linked to a

RNA (TR) component genes. The TR gene promoter can be linked to a

Cancer, especially neoplasias. The telomerase is necessary for the relation of cancer, especially neoplasias. The telomerase is necessary for the currestricted proliferative capacity of many human cancers. Mutation or dysregulation of the telomerase expression in cancer. Bubstances,

Con upregulation of telomerase expression in cancer. Substances,

clantified in the methods, can be used to block transcription from the TR gene promoter through interaction of the 5' regulatory sequences. These obstances, e.g. antisense oligonucleotides, transcription factors,

Competide nucleic acids and factors that disrupt signal transduction, are useful for cancer therapy. In particular, gene therapy vectors

Competide can be used to convert a promoter and a viral thymidine kinase gene can be used to convert a promoter and a viral thymidine compasia can be controlled or treated. Direct down-regulation of telomerase RNA gene through manipulation of transcription factors may be effective anticancer therapy and the cloning of the hTR gene promoter allows the analysis of therapeutic molecules which modulate hTR promoter activity. Sequences PAZ07324-332 and AAZ0032-340 represent constructs

With hTR promoter sequence element mutations
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                      Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR;
gene therapy; thymidine kinase gene; anticancer therapy; human; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; gene therapy; thymidine kinase gene; anticancer therapy; human; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mutated hTR promoter fragment containing construct 29112 (mSpl.4).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 51 BP; 8 A; 8 C; 21 G; 14 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                                                                                           (CANC-) CANCER RES CAMPAIGN TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Fig 19; 109pp; English
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les 48; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
Synthetic.
                                                                                              sapiens
                                                                                                                                                              WO9938964-A2
                                                                                                                                                                                                                                                            29-JAN-1999;
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                                                                                                                                                                                                           05-AUG-1999.
                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                          Keith WN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAZ00335;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mouse
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Matches
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The invention relates to promoter regions from mouse and human telomerase
RNA (TR) component genes. The TR gene promoter can be linked to a
Carcer, especially a gene encoding a cytotoxin, for therapy of
Carcer, especially neoplasias. The telomerase is necessary for the
Unrestricted proliferative capacity of many human cancers. Mutation or
dysregulation of the telomerase expression pathway may cause reactivation
Cor upregulation of telomerase expression in cancer. Substances,
conspiratores, e.g. antisense oligonucleotides, transcription from the TR
conspiratores, e.g. antisense oligonucleotides, transcription factors,
peptide nucleic acids and factors that disrupt signal transduction, are
useful for cancer therapy. In particular, gene therapy vectors
cospecially proffed-coddupp) comprising the promoter and a viral thymidine
kinase gene can be used to convert a productug, e.g. gancyclovir, so that
neoplasia can be controlled or treated. Direct down regulation of
telomerase RNA gene through manipulation of transcription factors may be
effective anticancer therapy and the cloning of the hTR gene promoter
allows the analysis of therapeutic molecules which modulate hTR promoter
allows the analysis of therapeutic molecules which modulate hTR promoter
activity.
                                                                                                                                                                                                                                                                                                                                                                                                                Mouse and human telomerase RNA gene promoters, useful for tumor specific
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    with hTR promoter sequence element mutations
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Pred. No. 2.6;
0; Mismatches
                                                                                                                                                          RES CAMPAIGN TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Fig 19; 109pp; English
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99WO-GB000308
                                                                             98GB-00001902
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                                                                                                                                                                                                                                                                                                                          WPI; 1999-479183/40.
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Best Local Similarity
Matches 48; Conserv
                                                                                                                                                          (CANC-) CANCER
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29-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               gene therapy
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Matches
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                                                                                                                                                                                            RNA (TR) component genes. The TR gene promoter can be linked to a heterologous gene, especially a gene encoding a cytotoxin, for therapy of cancer, especially neoplasias. The telomerase is necessary for the unrestricted proliferative capacity of many human cancers. Mutation or dysregulation of the telomerase repression pathway may cause reactivation or telomerase expression in cancer. Substances, identified in the methods, can be used to block transcription from the TR gene promoter through interaction of the 5' regulatory sequences. These substances, e.g. antisense oligomucleotides, transcription factors, peptide nucleic acids and factors that disrupt signal transduction, are useful for cancer therapy. In particular, gene therapy vectors (specially pGT62-codAupp) comprising the promoter and a viral thymidine kinase gene can be used to convert a prodrug, e.g. gancyclovir, so that neoplasia can be used to convert a prodrug, e.g. gancyclovir, so that the neoplasia can be used to convert a prodrug, e.g. gancyclovir, so that the neoplasia can be through manipulation of transcription factors may be telective anticancer therapy and the cloning of the hrR gene promoter allows the analysis of therapeutic molecules which modulate hrR promoter allows the analysis of therapeutic molecules which modulate hrR promoter activity. Sequences AAZ07324-332 and AAZ00332-340 represent constructs
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                                                                                                                                                                                  The invention relates to promoter regions from mouse and human telomerase
                                                                                                                   Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; gene therapy; thymidine kinase gene; anticancer therapy; human; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                       with hTR promoter sequence element mutations
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       8.8%; Score 39.6; DE
31.3%; Pred. No. 8.9;
ve 0; Mismatches
                                      (CANC-) CANCER RES CAMPAIGN TECHNOLOGY.
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                                                                                                                                                          Disclosure; Fig 19; 109pp; English
             98GB-00001902.
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les 42; Conservative
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                                                                                         WPI; 1999-479183/40
             29-JAN-1998;
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                                                                                                                                  gene therapy
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                                                                Keith WN
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LIVE INTURLION TEALEURS LO PERONCER FEGIONS FLOW mouse and numer LEGIONGEAGE

RNA (TR) component genes. The TR gene promoter can be linked to a carcer, especially neoplasias. The telomerase is necessary for the cancer, especially neoplasias. The telomerase is necessary for the currestricted proliferative capacity of many human cancers. Mutation or dysregulation of the telomerase expression pathway may cause reactivation or dysregulation of the telomerase expression in cancer. Substances, identified in the methods, can be used to block transcription from the TR came promoter through interaction of the 5' regulatory sequences. These substances, e.g. antisense oligomucleotides, transcription factors, peptide nucleic acids and factors that disrupt signal transduction, are useful for cancer therapy. In particular, gene therapy vectors (especially pGT62-codAupp) comprising the promoter and a viral thymidine cusquil for cancer therapy in particular, gene therapy vectors may be effective anticancer therapy and the clonding of the hTR gene promoter callows the analysis of therapeutic molecules which modulate hTR promoter callows the analysis of therapeutic molecules which modulate hTR promoter callows the hTR promoter sequence element mutations
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                                                                 Mouse and human telomerase RNA gene promoters, useful for tumor specific
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                                                                                                                                                                                                                                           invention relates to promoter regions from mouse and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
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Pred. No. 8.9;
0; Mismatches
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                                                                                                                                                                      Disclosure, Fig 19; 109pp; English
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WPI; 1999-479183/40.
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RESULT
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                          The invention relates to promoter regions from mouse and human telomerase
RNA (TR) component genes. The TR gene promoter can be linked to a
Reterologous gene, especially a gene encoding a cytotoxin, for therapy of
cancer, especially neoplasias. The telomerase is necessary for the
cancer, especially neoplasias. The telomerase is necessary for the
currestricted proliferative capacity of many human cancers. Mutation or
dysregulation of the telomerase repression pathway may cause reactivation
of upregulation of telomerase expression in cancer. Substances,
clantified in the methods, can be used to block transcription from the TR
gene promoter through interaction of the 5' regulatory sequences. These
substances, e.g. antisense oligonucleotides, transcription factors,
peptide nucleic acids and factors that disrupt signal transduction, are
useful for cancer therapy. In particular, gene therapy vectors
cuseful for cancer therapy. In particular, gene therapy vectors
chepcially pGF2-codAupp) comprising the promoter and a viral thymidine
kinase gene can be used to convert a producy, e.g. gancyclovir, so that
neoplasia can be controlled or treated. Direct down-regulation of
telomerase RNA gene through manipulation of transcription factors may be
effective anticancer therapy and the cloning of the hTR gene promoter
allows the analysis of therapeutic molecules which modulate hTR promoter
activity. Sequences AAZ07324-332 and AAZ00332-340 represent constructs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human telomerase RNA gene (hTR) promoter specific primer hll.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 GTGGAGGGTGGGCCTGGGTAAGGTAATGCCCATTTTTTGTCTAACC 47
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                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 47 BP; 8 A; 7 C; 18 G; 14 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                            with hir promoter sequence element mutations
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Disclosure; Fig 19; 109pp; English.
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Best Local
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cancer, especially neoplasias. The telomerase is necessary for the unrestricted proliferative capacity of many human cancers. Mutation or dysregulation of the telomerase repression pathway may cause reactivation or upregulation of telomerase expression in cancer. Substances, in dentified in the methods, can be used to block transcription from the TR gene promoter through interaction of the 5' regulatory sequences. These substances, e.g. antisense oligonucleotides, transcription factors, useful for cancer therapy. In particular, gene therapy vectors (especially pGT62-codAupp) comprising the promoter and a viral thymidine kinase gene can be used to convert a product down-regulation of the neoplasia can be used to convert a product down-regulation of telomerase RNA gene through manipulation of transcription factors may be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to promoter regions from mouse and human telomerase RNA (TR) component genes. The TR gene promoter can be linked to a heterologous gene, especially a gene encoding a cytotoxin, for therapy of cancer, especially neoplasias. The telomerase is necessary for the unrestricted proliferative capacity of many human cancers. Mutation or dysregulation of the telomerase repression pathway may cause reactivation or upregulation of telomerase expression in cancer. Substances, identified in the methods, can be used to block transcription from the TR
                                                                                                                                                                                                                                                                                                            effective anticancer therapy and the cloning of the hTR gene promoter allows the analysis of therapeutic molecules which modulate hTR promoter activity. Sequences AAZ07696-321 represent PCR primers used in cloning
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14;
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                                                                                                                                                                                                                                                                                                                                                                                  and mutagenisis of human TR gene (hTR) promoter region
                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 38 BP; 2 A; 7 C; 23 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GGGTTGCGGAGGGTGGCCTGGGAGGGGTGGTGGCC 38
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Matches 36; Conservative
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                   substances, e.g. antisense oligonucleotides, transcription factors, peptide nucleic acids and factors that disrupt signal transduction, are useful for cancer therapy. In particular, gene therapy vectors are (especially pGT62-codAup) comprising the promoter and a viral thymidine kinase gene can be used to convert a prodrug, e.g. gancyclovir, so that neoplasia can be controlled or treated. Direct down-regulation of telomerase RNA gene through manipulation of transcription factors may be effective anticancer therapy and the cloning of the hTR gene promoter allows the analysis of therapeutic molecules which modulate hTR promoter activity. Sequences AZGV769-321 represent PCR primers used in cloning and mutagenisis of human TR gene (hTR) promoter region
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                                                                                                                                                                                                                                                                                                                                                             Gaps
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promoter through interaction of the 5' regulatory
                                                                                                                                                                                                                                                                    Sequence 38 BP; 3 A; 7 C; 21 G; 7 T; 0 U; 0 Other;
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Pred. No. 36;
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Best Local Similarity 91.7%;
Matches 33; Conservative
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kinase gene can be used to convert a prodrug, e.g. gancyclovir, so that neoplasia can be controlled or treated. Direct down-regulation of telomerase RNA gene through manipulation of transcription factors may be effective anticancer therapy and the cloning of the hTR gene promoter allows the analysis of therapeutic molecules which modulate hTR promoter activity. Sequences AA207669-321 represent PCR primers used in cloning and mutagenisis of human TR gene (hTR) promoter region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; telomerase RNA component; anticancer therapy; purification; assay; vaccine; cancer; antisense oligonuclectide; ss.
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                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                               Antisense oligonucleotide anti-P for human telomerase RNA component.
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                                                                                                                                                                DB 1; Length 38;
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                                                                                                                                                                                                Indels
                                                                                                                                Sequence 38 BP; 5 A; 7 C; 20 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                               36
                                                                                                                                                                                                                                                              3 gegingcegaaaareeeccreeeaeeereereecc 38
                                                                                                                                                                                                                               1 GGGTTGCCGAGGGTGGGCCTGGGAGGGGTGGTGGCC
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                                                                                                                                                              Score 31.2; DE Pred. No. 36; 0; Mismatches
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"biotinylated"
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"biotinylated"
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                                                                                                                                                                6.98;
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ID AAV63645 standard; DNA; 31
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Best Local Similarity
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biochemical analysis of its activity, and in preparation of antibodies. Fragments of telomerase, or nucleic acid encoding them, are used in vaccines, and for treating over expression of telomerase, particularly in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New peptide nucleic acid (PNA) compounds that inhibit telomerase activity in mammalian cells is useful as probes to detect the RNA component of a mammalian telomerase.
                                                                                                                                                                                                                                                                                                                                                          Peptide nucleic acid; PNA; telomerase; ribonucleoprotein enzyme; cancer; inhibitor; neoplasia; neurodegenerative disease; aging; hyperplasia; AIDS; HIV; fungal infection; forensic identification; detect; tumour; paternity testing; ss.
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                                                                                             Query Match 6.9%; Score 31; DB 1; Length 31; Best Local Similarity 100.0%; Pred. No. 28; Matches 31; Conservative 0; Mismatches 0; Indels
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                                                                    Sequence 31 BP; 7 A; 9 C; 7 G; 8 T; 0 U; 0 Other;
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                                                                                                                                                     42 TIGICTAACCCTAACTGAGAAGGGCGTAGGC 72
                                                                                                                                                                     31 TTGTCTAACCGTAAGGGGGGGGGCGTAGGC 1
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                                                                                                                                                                                                                                                                                                                                         Telomerase target sequence.
                                                                                                                                                                                                                                                     AAA37591 standard; RNA; 31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to peptide nucleic acids (PNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in mammalian cells by hybridising to the RNA component of mammalian telomerase. The PNAs are useful as probes to detect the RNA component of
probes to detect or quantitate polynucleotides having a human telomerase RNA (hTR) sequence. PNA probes are also used for forensic identification of individuals. e.g. paternity testing, based on hTR gene restriction fragment length polymorphism (RFLP) pattern. PNAs are also useful as probes to detect the RNA component of a mammalian telomerase and as inhibitors of telomerase activity. The method of the present invention possibly at an earlier stage, before cells are detected as cancerous based on pathological characteristics. The diagnostic and prognostic methods of the present invention can be used to detect an immortal or neoplastic cell or tumour tissue or cancer of any origin, provided the cell expresses telomerase activity and its RNA component
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6.9%; Score 31; DB 1; Length 31; 71.0%; Pred. No. 28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
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/*tag= a
/note= "Telomerase RNA active site"
                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 31 BP; 8 A; 6 C; 8 G; 0 T; 9 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       40 TITIGICTAACCCTAACTGAGAAGGGCGTAG 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Corey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      9; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human telomerase RNA (hTR) strand #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 2; Col 37-38; 46pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-638024/73.
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Best Local Similarity
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09-APR-1997;
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WPI; 2003-465598/44.
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04-APR-1997;
18-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          modified base
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                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                            RESULT 18
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          detect and/or quantitate polynucleotide having the human telomerase RNA component (hTR) sequence, as well as in forensic identification of criminal individuals, such as paternity teating or identification of criminal suspects or unknown descendants based on the hTR gene RFLP pattern. The PNA can be further used for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative alseases. The PNAs in combination with other pharmaceuticals (such as antinoeplastic or cytostatic agents) can be used for treating neoplasia, hyperplasia, human immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomera metabolism or telomerase activity. The present sequence representing HTR RNA strand #2 is used to test the inhibition of telomerase activity by the PNAs of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Purified human telomerase, useful for inducing immune response in animals, comprises several thousand folds increased purity compared with cytoplasmic crude cell preparations.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lichtsteiner SP, Vasserot AP, Pruzan RA;
                                                                                                                                                                                                                                          Gaps
 mammalian telomerase and as inhibitors of telomerase activity, or to
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Human, Telomerase, vaccine, antibody, cancer, EF2H, nucleolin, antisense oligonucleotide, anti-P, ss.
                                                                                                                                                                                                               Score 31; DB 1; Length 31;
Pred. No. 28;
9; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                        Antisense oligonucleotide for human telomerase, anti-P.
                                                                                                                                                                                          Sequence 31 BP; 8 A; 6 C; 8 G; 0 T; 9 U; 0 Other;
                                                                                                                                                                                                                                                                40 TTTTGTCTAACCCTAACTGAGAAGGGCGTAG 70
                                                                                                                                                                                                                                                                               UNUUGUCUAACCCUAACUGAGAAGGGCGUAG
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/note= "G is biotinylated"
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                                                                                                                                                                                                                                Local Similarity 71.0 tes 22; Conservative
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04-APR-1997;
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                                                                                                                                                                       invention
                                                                                                                                                                                                                                                                                                                                                                           AAS09472;
                                                                                                                                                                                                                   Query Match
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component has DNA polymerase activity and a molecular weight of 200-2000 component has DNA polymerase activity and a molecular weight of 200-2000 kilo Daltons (kDa). The purified telomerase is useful for inducing a chamerase or immunogenic fragments are useful as useful for inducing a chamerase or immunogenic fragments are useful as vaccines for treating diseases associated with over-expression of telomerase, such as cancer and for producing antibodies that recognize telomerase, which are useful as a ffinity agents in isolating the proteins and for detecting the presence of proteins in a sample, such as call or tissue. Identification of telomerase aids in diagnosis of cancer or pre-cancerous states.

Telomerase and/or telomerase associated proteins are also useful for screening compounds to identify agents that alter the association of telomerase associated proteins are also useful for screening compounds to identify agents that alter the association of telomerase associated proteins or BF2H with telomerase
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6.9%; Score 31; DB 1; Length 31; 00.0%; Pred. No. 28;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human telomerase antisense oligonucleotide primer AntiP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 31 BP; 7 A; 9 C; 7 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Telomerase; antisense; primer; AntiP; ss; cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              42 TIGICIAACCCTAACTGAGAAGGGCGTAGGC 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Pred. No. 28;
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"Biotinylated"
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/note= "Biotinylated"
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97US-00833377.
99US-00420056.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABX10982 standard; DNA; 31
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Matches 31; Conservative
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/note= '
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measuring telomerase enzymatic activity of the composition in presence of a regulator. The telomerase protein of the invention may be used in developing and testing assays for measuring telomerase activity which are useful in characterising cancer and pre-cancer cells, for identifying and testing regulators of telomerase activity in in vitro assay and for preparing antibodies against telomerase. The mammalian telomerase protein of the invention is at least approximately 3000 fold more pure (in terms of telomerase activity per weight of protein) than a crude extract of cell from adenovirus- transformed kidney cell. Purified telomerase daralizers a thorough blochemical analysis of the anzyme's mechanism for developing mechanism- based regulators. The present sequence represents a human telomerase antisense oligonucleotide which has affinity to telomerase and is used to purify the telomerase protein of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 31; DB 1; Length 31;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 31 BP; 7 A; 9 C; 7 G; 8 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TTGTCTAACCCTAACTGAGAAGGGCGTAGGC 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Pred. ...
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-NOV-2000; 2000US-00717829
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6.98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADC35648 standard; DNA; 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   31; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-742824/70.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 JS6545133-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-AUG-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-APR-1997;
18-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       38-APR-2003
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IID ADC33

AAC ADC34

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containing telomerase protein with an anion exchange matrix, and collecting protein that binds the matrix, combining a fraction containing telomerase protein with a cation exchange matrix (such as a heparin matrix), and collecting protein that binds the matrix. The method
                                                        comprises successively enriching fractions containing telomerase protein on several different ion exchange matrices and combining a fraction containing telomerase protein with an intermediate selectivity matrix, collecting protein that binds the matrix, where the intermediate selectivity matrix and separating a fraction containing the telomerase protein by gel filtration chromatography or gradient centrification. The telomerase is enriched from an extract of cells stably expressing telomerase. This sequence represents an antisense oligonucleotide to the
                                                                                                                                                                telomerase. This sequence represents an antisense oligonucleotidé to t}
RNA component of human telomerase that can be used in the purification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pruzan RA;
                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Telomerase activity; therapy; cancer; cytostatic; antisense; ss.
                                                                                                                                                                                                                                                                                       ô
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Identifying telomerase regulators useful for treating cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Vasserot AP,
                                                                                                                                                                                                                                                       DB 1; Length 31;
28;
                                                                                                                                                                                                                                                                                     Indele
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human telomerase RNA antisense oligonucleotide, anti-P.
                                                                                                                                                                                                                           Seguence 31 BP; 7 A; 9 C; 7 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                       ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lichtsteiner SP,
                                                                                                                                                                                                                                                                                                                   42 TTGTCTAACCCTAACTGAGAAGGGCGTAGGC 72
                                                                                                                                                                                                                                                                       100.0%; Pred. No. 28; ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                rrercraacceraacreagaagggggggg 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /mod_base= OTHER
/note= "Biotin labelled"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /mod_base= OTHER
/note= "Biotin labelled'
                                                                                                                                                                                                                                                           6.9%; Score 31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /*tag= b
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                           BP
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04-APR-1997; 97US-00833377.
18-OCT-1999; 99US-00420056.
20-NOV-2000; 2000US-00717828.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-DEC-2002; 2002US-00330872
                                                                                                                                                                                                                                                                                                                                                                                                                           ADG62870 standard; DNA; 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11-MAR-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /*tag= a
                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 100.
Matches 31; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Δ.
                                                                                                                                                                                                 nethod of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Weinrich SL, Atkinson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WEINRICH S L.
ATKINSON E M.
LICHTSTEINER S
VASSEROT A P.
PRUZAN R A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                VPI; 2003-811733/76.
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         02-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADG62870;
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(LICH/)
(VASS/)
(PRUZ/)
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                                                                                                                                                                                                                                                                                                                                                                                              RESULT 20
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                            The invention relates to a method for identifying regulators of telomerase activity that may be useful for treating cancers. The method may be used to identify regulators e.g. antibodies, of telomerase activity which may be useful as cancer treatments. It has been found that found that the cells of many human cancers have telomerase activity. This helps explain why cancer cells continue dividing without becoming senescent. If telomerase activity in cancer cells can be inhibited, the cancer cells are expected to reach senescence and cease dividing. The present sequence is human telomerase antisense oligonulceotide used to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mouse, rat and chinese hamster are all claimed. Antisense deligiouscleotides can be used to block the activity of the telomerase; probes and primers can be used in detection; vectors and host cells transformed with the isolated telomerase genes can be used for production of telomerases; RNA and DNA ribozymes and triplex forming oligonucleotides directed against the telomerase genes can be used therapeutically as can plasmids. A mouse which lacks the telomerase gene (also claimed) can be used for study of telomere regulation in vivo, and the role it plays in immortalisation. Three primers (AAT11040, AAT11043,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    - useful in studying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The RNA components of (non) human mammalian telomerase(s) especially from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Telomerase; mammal; antisense; triplex forming oligonucleotide; plasmid;
                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Marhuenda MA;
                                                                                                                                                                                                                                                                           ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Primer for production of telomerase antisense oligonucleotide.
                                                                                                                                                                                                                                         DB 1; Length 31; 28;
                                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Greider C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RNA components of (non)human mammalian telomerase(s) cell senescence and immortalisation.
                                                                                                                                                                                                        Sequence 31 BP; 7 A; 9 C; 7 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                    42 TTGTCTAACCCTAACTGAGAAGGGCGTAGGC 72
                                                                                                                                                                                                                                                                                                                        31 TTGTCTAACCCTAACTGAGAAGGGCGTAGGC 1
                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Funk W,
                                                                                                                                                                          illustrate the method of the invention.
                                                                                                                                                                                                                                        6.9%; Score 31;
100.0%; Pred. No.
Disclosure; SEQ ID NO 1; 22pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Feng J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Page 53; 85pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COLD SPRING HARBOR LAB
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95US-00387524.
95US-00485778.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      94US-00272102.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        95WO-US008620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             probe; primer; ribozyme; ss
                                                                                                                                                                                                                                                                                                                                                                                                   043/c
AAT11043 standard; DNA; 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                        Query Match
Best Local Similarity 100.
Matches 31; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Avillon AA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1996-097428/10.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Villeponteau B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO9601614-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                02-JUL-1996
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13-FEB-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-JUN-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-JAN-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAT11043;
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AAT11044) were used to produce antisense oligonucleotides which were then used to produce antisense expression plasmids. AAT11040 was used alongside both AAT11043 and AAT11044 to produce two different antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence is a PCR primer for a RNA component of human library obtd. From the human ling fibroblast cell line Will. 38. The RCHT can be used in the recombinant prodn. of an active telomerase mol. capable of adding sequences to chromosomal DNA telomeres, and in the resumbinant prodn. of an active telomerase mol. capable of adding sequences for the detection of mutant mammalian telomerase RNA component polynuclectides. The RCHT may also be used in the identification of telomerase modulating agents, and in the detection of telomerase related, or neoplastic conditions in a patient. Polynuclectides of at least 25 consecutive nucleotides identical, or complementary to the RFHT sequence linked to heterologous transcriptional regulatory sequences, can be used for the gene therapy of human diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         esp. human - useful in identifying
                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RNA component; human; telomerase; lung fibroblast; cell line WI-38;
                                                                                                                                                                                                                                                                                                                                                                                                                                        RNA component of human telomerase antisense plasmid PCR primer Gl.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            recombinant production; synthesis; mutant; detection; mammalian; identification; modulating agent; neoplastic condition; transcriptional regulatory sequence; gene therapy; disease; polymerase chain reaction; antisense plasmid; PCR primer; ss.
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0
                                                                                                                            6.7%; Score 30; DB 1; Length 30;
100.0%; Pred. No. 32;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1; Length 30; 32;
                                                                                        Sequence 30 BP; 13 A; 6 C; 11 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 30 BP; 13 A; 6 C; 11 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Funk W, Andrews WH;
                                                                                                                                                                                                  77 IGCTTTTGCTCCCCCCCCCCTGTTTTCTC 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          candidate telomerase-modulating agents.
                                                                                                                                                                                                                                    30 recrirricarcacacacacacitriria 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RNA component of mammalian telomerase,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 8; Page 80; 114pp; English
                                                                                                                                                                                                                                                                                                                                  BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 94US-00272102.
94US-00330123.
95US-00472802.
95US-00482115.
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                                                                                                       Query Match
Best Local Similarity 100...
Conservative
                                                                                                                                                                                                                                                                                                                               AAT10298 standard; DNA; 30
                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Villeponteau B, Feng J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1996-097581/10.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO9601835-A1
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                                                       molecules
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                   AAT10298;
                                                                                                                                                                                                                                                                                              RESULT 22
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Human; telomerase RNA component; anticancer therapy; purification; assay; vaccine; cancer; antisense oligonuclectide; ss.
                                                                                                      Antisense oligonucleotide 14 for human telomerase RNA component.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Purification of telomerase on affinity material - useful for,
                                                                                                                                                                                                                                                                                                                                                                                                                                                   diagnosis and treatment of cancer.
                                 AAV63649 standard; DNA; 30 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1998-594485/50.
                                                                                                                                                                                                                                                                                                                                                       (GERO-) GERON
                                                                                                                                                                                                  Key
modified_base
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                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                            WO9845450-A1
                                                                                                                                                                                                                                                                                                         04-APR-1997;
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                                                                               15-FEB-1999
                                                                                                                                                                                                                                                                                   15-OCT-1998
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                                                                                                                                                                Synthetic.
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                                                        AAV63649;
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                   AAV63649/c
ID AAV636
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present sequence represents an antisense oligonucleotide directed against the human telomerase RNA component gene sequences. The oligonucleotide can be used as an affinity agent in the methods of the invention, which are used to purify human telomerase. The methods of the use of several sequential steps, including the use of two matrixes that bind molecules bearing negative charges, a matrix that binds molecules bearing negative charges, an matrix that binds molecules bearing negative charges, an affinity purification step and a size separation. Telomerase is a particular target of anticancer therapies, and is useful in assays for characterizing (pre)cancerous cells. Telomerase can also be used to screen for specific modulators, for biochemical analysis of its activity, and in preparation of antibodies. Fragments of telomerase, or nucleic acid encoding them, are used in vaccines, and for treating over expression of telomerase, particularly in
                                                                                                                                                                                                Human; telomerase RNA component; anticancer therapy; purification; assay;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Atkinson EM, Lichtsteiner SP, Vasserot AP, Pruzan RA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Purification of telomerase on affinity material - useful for, e.g. diagnosis and treatment of cancer.
                                                                                                                                                                           Antisense oligonucleotide 13 for human telomerase RNA component.
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 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 30 BP; 6 A; 6 C; 8 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                             vaccine; cancer; antisense oligonucleotide; ss.
                       TGCTTTTGCTCCCCGCGCGCTGTTTTCTC 106
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                    30 TGCTTTTGCTCCCCGCGCCCTGTTTTTCTC 1
                                                                                                                                                                                                                                                                                              /*tag= a
/note= "biotinylated"
                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 24; 76pp; English.
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                                                                                                      В
                                                                                                                                                                                                                                                                                                                                                                               97WO-US006012.
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                                                                                                      AAV63648 standard; DNA; 30
                                                                                                                                                    (first entry)
 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1998-594485/50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                             CORP.
                                                                                                                                                                                                                                                                        Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                             (GERO-) GERON
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Weinrich SL,
                                                                                                                                                                                                                                                                                                                                  WO9845450-A1
                                                                                                                                                                                                                                                 Ното варіепв
                                                                                                                                                                                                                                                                                                                                                                               04-APR-1997;
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                                                                                                                                                    15-FEB-1999
 30;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kealey JT;
                                                                                                                                                                                                                                     Synthetic.
                                                                                                                              AAV63648;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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                                                                               Best Loc
 Matches
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Pruzan RA;

Vasserot AP,

Lichtsteiner SP,

Atkinson EM,

/*tag= a /note= "biotinylated" Location/Qualifiers

(first entry)

97WO-US006012. 97WO-US006012

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                                                      The present sequence represents an antisense oligonucleotide directed against the human telomerase RNA component gene sequences. The oligonucleotide can be used as an affinity agent in the methods of the invention, which are used to purify human telomerase. The methods of the the use of several sequential steps, including the use of two matrixes that bind molecules bearing negative charges, a matrix that binds molecules bearing negative charges, a matrix that binds size separation. Telomerase is a particular target of anticancer therapies, and is useful in assays for characterizing (pre) cancerous cells. Telomerase can also be used to screen for specific modulators, for biochemical analysis of its activity, and in preparation of antibodies. Fragments of telomerase, or nucleic acid encoding them, are used in vaccines, and for treating over expression of telomerase, particularly in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1; Length 30; 32;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 30 BP; 8 A; 5 C; 13 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6.7%; Scor.
100.0%; Pred. No. Jr.
7. 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         137 CCTGCCGCCTTCCACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cerececerrecacerrearreragase 1
Disclosure; Page 24; 76pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
nes 30; Conserv
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AAV41175/c
ID AAV41
XX
AC AAV41
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AAACAAAAAATGTCAGCTGCTGGCCGTTC 196

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AAACAAAAATGTCAGCTGCTGGCCCGTTC 1

RNA component; human telomerase; antisense oligonucleotide; infection; neuroblastoma; bladder cancer; colon cancer; prostate cancer; cancer; contraception; sterilisation; immunosuppression; therapeutic; hTR; immune system down-regulation; anti-inflammatory therapy; ss.

RNA component of human telomerase (hTR) antisense oligo 16.

08-OCT-1998 (first entry)

US09436060A-16.rng.sl

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RNA component; human telomerase; antisense oligonucleotide; infection; neuroblastoma; bladder cancer; colon cancer; prostate cancer; cancer; contraception; sterilisation; immunosuppression; therapeutic; hTR; immune system down-regulation; anti-inflammatory therapy; ss.
                     RNA component of human telomerase (hTR) antisense oligo 21.
                                                                                                                                                                                  Kim NW, Wu F, Kealey JT, Pruzan R, Weinrich SL;
                                                                                                                                97WO-US023619.
                                                                                                                                             96US-00770564.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAV41172 standard; DNA; 30 BP.
        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                              30; Conservative
                                                                                                                                                                                                  WPI; 1998-377670/32.
                                                                                                                                                                    (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                   Homo sapiens,
                                                                                                 WO9828442-A1
                                                                                                                               19-DEC-1997;
                                                                                                                                                     20-DEC-1996;
        08-OCT-1998
                                                                                                                                              20-DEC-1996;
                                                                                                               02-JUL-1998.
                                                                           Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAV41172/
ID AAV4
XX
AC AAV4
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New polynucleotide(s) anti:sense to human telomerase - used for detecting or inhibiting human telomerase, e.g. for treating cancers, contraception, immuno-suppression or treating infection.

Claim 11; Page 65; 80pp; English.

Pruzan R, Weinrich SL;

Kim NW, Wu F, Kealey JT,

(GERO-) GERON CORP.

WPI; 1998-377670/32.

97WO-US023619. 96US-00770564. 96US-00770565.

19-DEC-1997; 20-DEC-1996; 20-DEC-1996;

02-JUL-1998

WO9828442-A1. Homo sapiens

Synthetic

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Sequences shown in AAV41169 to AAV41181 represent antisense
oligonucleotides to the RNA component of human telomerase (hTR). These
antisense oligonucleotides specifically hybridise to a nucleotide
sequence within an accessible region of the HTR, but that does not
bybridise to a sequence within the template region of hTR. These
component of human telomerase in a sample. This is useful for diagnosing
component of human telomerase in a sample. This is useful for diagnosing
cancer (especially neuroblastoma, bladder, colon and prostate cancer),
component of human telomerase in a sample. The inhibitory
cancer (especially neuroblastoma, bladder, colon and prostate cancer),
coligonucleotides can inhibit the telomerase activity level in a cell by
coligonucleotides can inhibit the telomerase activity level in a cell by
coligonucleotides and inhibit the telomerase activity inhibiting the half-
life of the telomerase RNA component transcript, inhibiting the half-
coligonoment into the telomerase holoenzyme, or inhibiting the
polymerase activity of telomerase activity in both cultured cells and in
preventing cancer, for contraception or sterilisation, for
calls in vivo. They can be used in therapeutics for treating or
preventing cancer, for contraception or sterilisation, for
immunosuppression, and for selectively down-regulating specific branches
colimnatory therapies or for treating infections by, e.g. yeast,
New polynucleotide(s) anti:sense to human telomerase - used for detecting or inhibiting human telomerase, e.g. for treating cancers, contraception, immuno-suppression or treating infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 30 BP; 8 A; 5 C; 13 G; 4 T; 0 U; 0 Other;
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AAZ23627 standard; DNA; 30 BP
Query Match
Best Local Similarity 100.
Matches 30; Conservative
                                                                               30
                                                                                                                                  AAZ23627/
ID AAZ2
                                                                                                                     RESULT 27
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                                       Gaps
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0
           1; Length 30;
                                    0; Indels
                                                           137 CCIGCCGCCTTCCACCGTTCATTCTAGAGC 166
       6.7%; Score 30; DB
100.0%; Pred. No. 32;
ilve. 0; Mismatches
                                                                          CTGCCGCCTTCCACCGTTCATTCTAGAGC
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Sequences shown in AAV41169 to AAV41181 represent antisense
cligonuclectides to the RNA component of human telomerase (hTR). These
antisense oligonuclectides specifically hybridise to a nuclectide
capture within an accessible region of the hTR, but that does not
component of human telomerase in a sample. This is useful for diagnosing
cancer (especially neuroblastoma, bladder, colon and prostate cancer),
component of human telomerase in a sample. This is useful for diagnosing
cancer (especially neuroblastoma, bladder, colon and prostate cancer),
cancer (especially neuroblastoma, che minibitory
cancer (especially neuroblastoma, concerning the half-
call controlled component transcript, inhibiting assembly of
the RNA component into the telomerase holoenzyme, or inhibiting telomerase activity in both cultured cells and in
calls in vivo. They can be used in therapeutics for treating or
cancer inhibiting telomerase activity in both cultured cells and in
calls in vivo. They can be used in therapeutics for treating or
cancer inflammatory therapeuts or for treating infections by, e.g. yeast,
cancer inflammatory theraples or for treating infections by, e.g. yeast,
cancer inference or paragraphs or for treating infections by, e.g. yeast,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 30 BP; 6 A; 10 C; 9 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  290 CTGCCACCGCGAAGAGTTGGGCTCTGTCAG 319
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Pred. No.
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Kealey JT;

Pruzan RA,

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This invention describes a novel composition comprising human telomerase having at least 2000-fold (preferably at least 6000-fold) increased relative purity compared with crude extract of cells from adenovirus-transformed kidney cell line. The composition is useful for eliciting an immune response in animals and may therefore be used as a vaccine for reacting diseases associated with the overexpression of telomerase e.g. cancer. AAZ23626-Z23637 represent oligonucleotides used in the isolation of human clone 28-1 which contains a fragment of the human telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             human; immune response; cancer; vaccine; treatment; disease;
                                                                                                                                                                                                                                                                                                                   useful for treating diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1; Length 30; 32;
                                                                                                                                                                                                                                                                                                                 Compositions comprising human telomerase, useful for tree associated with overexpression of telomerase e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human clone 28-1 telomerase oligonucleotide oligo-14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Seguence 30 BP; 6 A; 6 C; 8 G; 10 T; 0 U; 0 Other;
   /*tag= a
/note= "5'-biotinylated guanosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /*tag= a
/note= "5'-biotinylated guanosine"
                                                                                                                                                                                                                               Weinrich SL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             167 AAACAAAAATGTCAGCTGCTGGCCGTTC 196
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 6.7%; Score 30; DB Best Local Similarity 100.0%; Pred. No. 32; Matches 30; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30 AAACAAAAATGTCAGCTGCTGGCCCGTTC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Col 45-46; 34pp; English.
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                                                                                                                                                                                                                               Lichtsteiner SP,
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ID AAZ23631 Btandard; DNA; 30
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/*tag=
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modified_base
                                                                                                                        04-APR-1997;
                                                                                                                                                           04-AUG-1995;
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                                                                                                                                                                                                                             Atkinson EM,
Vasserot AP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-JAN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Telomerase;
                                                     JS5968506-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US5968506-A
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                                                                                     19-OCT-1999
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                                                                                                                                                                                                                                                   Jasserot
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention describes a novel composition comprising human telomerase relative purity compared with created relative purity compared with created extract of cells from adenovizustransformed kidney cell line. The composition is useful for eliciting an immune response in animals and may therefore be used as a vaccine for treating diseases associated with the overexpression of telomerase e.g. cancer. AAZ23626.223637 represent oligomucleotides used in the isolation of human clone 28-1 which contains a fragment of the human telomerase
                                                     Telomerase; human; immune response; cancer; vaccine; treatment; disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Telomerase; human; immune response; cancer; vaccine; treatment; disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Compositions comprising human telomerase, useful for treating diseases associated with overexpression of telomerase e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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32;
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                    Human clone 28-1 telomerase oligonucleotide anti-P
                                                                                                                                                                                              /*tag= a
/note= "5'-biotinylated guanosine"
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                                                                                                                                                                                                                                                                                                                                                                                                                             Lichtsteiner SP, Weinrich SL,
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100.0%; Pred. No. ...
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           43 TGTCTAACCCTAACTGAGAAGGGCGTAGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Col 43-44; 34pp; English.
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                                                                                                                                                           Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                    97US-00833377
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Best Local Similarity 100.
Matches 30; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-590379/50.
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                                                                                                                                                           Key
modified_base
                                                                                                                                                                                                                                                                                                                      04-APR-1997;
                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                         04-AUG-1995;
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Vasserot AP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-JAN-2000
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                                                                         primer; ss
                                                                                                        Synthetic
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RESULT 28 AAZ23630/c

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Gaps

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The invention relates to polynucleotide inhibitors (I) and methods for inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity and proliferation of a telomerase positive cell, and in manufacturing a medicament for inhibiting telomerase activity in a cell and in treating a medicament for inhibiting telomerase activity in a cell cadenocarcinoma of breast, prostate or colon, mixed cell leukaemis. Consection of the construction of telomerase activity in a cell and in treating a tumour or in manufacturing a medicament for the cuseful in treating a tumour or in manufacturing a medicament for the consection of tumour. The polynucleotide inhibitors may also be used in diagnostic assays for detecting RNA or DNA. Inhibition of telomerase activity in cells in vivo is useful in prophylactic and therapeutic activity in cells in vivo is useful in prophylactic and therapeutic context in a concer and other disorders involving inappropriate expression of telomerase, and in treating veterinary proliferative chases. Inhibition of telomerase in haematopoietic stem cells is useful context in the cells in the context of the c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               branches of the immune system. The present sequence represents human telomerase polynucleotide inhibitor \#9, as described in the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Purified human telomerase, useful for inducing immune response in animals, comprises several thousand folds increased purity compared with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human, Telomerase, vaccine, antibody, cancer; BF2H, nucleolin; antisense oligonucleotide; Oligo 14; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      6.7%; Score 30; DB 1; Length 30;
100.0%; Pred. No. 32;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antisense oligonucleotide for human telomerase, Oligo 14.
tumors, Hodgkin's disease, or inflammatory conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 30 BP; 8 A; 5 C; 13 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Lichtsteiner SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              137 CCTGCCGCCTTCCACCGTTCATCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mod_base= G
/note= "G is biotinylated"
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                                                       3; Page 32; 48pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       95US-00510736.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Weinrich SL, Atkinson EM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /*tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity 100.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-450477/48.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             invention
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Matches
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AAS09476/c
THE STATE OF THE S
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                                                                                                                                                                                                                                                                                                                                                   This invention describes a novel composition comprising human telomerase having at least 2000-fold (preferably at least 6000-fold) increased relative purity compared with crude extract of cells from adenovirus-transformed kidney cell line. The composition is useful for eliciting an immune response in animals and may therefore be used as a vaccine for treating diseases associated with the overexpression of telomerase e.g. cancer. AAZ23626-Z23637 represent oligonucleotides used in the isolation of human clone 28-1 which contains a fragment of the human telomerase described in the method of the invention
                                                                                                                                                                                                        Compositions comprising human telomerase, useful for treating diseases associated with overexpression of telomerase e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma;
breast; prostate; colon; mixed cell leukaemia; Hodgkin's disease;
fertility; inflammatory condition; tumour; cancer; veterinary;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                          Kealey JT;
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                                                          Pruzan RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 30; DB 1; Length 30;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 30 BP; 8 A; 5 C; 13 G; 4 T; 0 U; 0 Other;
                                                          Lichtsteiner SP, Weinrich SL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human telomerase polynucleotide inhibitor #9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immunosuppression; telomerase inhibitor; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Pred. No. 32; cive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      137 CCTGCCGCCTTCCACCGTTCATTCTAGAGC 166
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                                                                                                                                                                                                                                                                                            Disclosure; Col 45-46; 34pp; English.
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                                                                                                                                                 WPI; 1999-590379/50
(GERO-) GERON CORP.
                                                          AP;
                                                          Atkinson
                                                                                         Vasserot
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Pruzan RA;

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17-JAN-2002
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ABA91517/c
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                                                                           The sequence represents a biotinylated antisense oligonucleotide used in the purification of human telomerase. The invention relates to a purified human telomerase core enzyme protein comprising 2000-fold increased burity compared with a crude extract of calls from adenovirus-transformed purity compared with a crude extract of calls from adenovirus-transformed component has DNA polymerase activity and a molecular weight of 200-2000 kilo baltons (kDa). The purified telomerase is useful for inducing a humoral or cell-mediated immune response in an animal. Purified telomerase or immunogenic fragments are useful as vaccines for treating diseases associated with over-expression of telomerase, which are useful as a fifnity agents in isolating the proteins and for detecting the proteins of proteins in a sample, such as cell or tissue. Identification of telomerase aids in disgnosis of cancer or pre-cancerous states.

Telomerase and/or telomerase associated proteins are also useful for screening compounds to identify agents that alter the association of telomerase.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Purified human telomerase, useful for inducing immune response in animals, comprises several thousand folds increased purity compared with cytoplasmic crude cell preparations.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pruzan RA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human, Telomerase, vaccine, antibody, cancer; EF2H, nucleolin; antisense oligonucleotide; Oligo 13; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 30; DB 1; Length 30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antisense oligonucleotide for human telomerase, Oligo 13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 30 BP; 8 A; 5 C; 13 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Pred. No. 32; ive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /mod_base= G
/note= "G is biotinylated"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30 ccreccecrrccaccerrcarrcragaec 1
cytoplasmic crude cell preparations.
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                                          Disclosure; Col 18; 29pp; English.
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AAS09475 standard; DNA; 30 BP.
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mod_base= G
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97US-00833377
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
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Best Local Similarity 100.
Matches 30; Conservative
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Kealey JT;
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ID AASO
AC AASO
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XX AASO
XX Huma.
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                                                   The sequence represents a biotinylated antisense oligonucleotide used in the purification of human telomerase. The invention relates to a purified human telomerase core enzyme protein comprising 2000-fold increased purity compared with a crude extract of cells from adenovirus-transformed widney cell line (293 cells) and when associated with telomerase RNA component has DNA polymerase activity and a molecular weight of 200-2000 kilo Daltons (kDa). The purified telomerase is useful for inducing a humoral or cell-mediated immune response in an animal. Purified a telomerase or immunogenic fragments are useful as vaccines for treating diseases associated with over-expression of telomerase, such as cancer and for producing antibodies that recognize telomerase, which are useful
                                                                                                                                                                                                                                                                                                                                                                                                                                               as affinity agents in isolating the proteins and for detecting the presence of proteins in a sample, such as cell or tissue. Identification of telomerase aids in diagnosis of cancer or pre-cancerous states. Telomerase and/or telomerase associated proteins are also useful for screening compounds to identify agents that alter the association of telomerase-associated proteins, such as nucleolin or EF2H with telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Amplifying signal by contacting nanoparticles (N) having bound oligonuclectides to silver deposited on substrate to form N-silver sandwich that is contacted with silver, reducing agent to form silver-N-silver sandwich.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Silver staining, signal detection; nanoparticle; diagnosis; screening;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oligonucleotide used in gold nanoparticle complex.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             167 AAACAAAAATGTCAGCTGCTGGCCCGTTC 196
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAACAAAAATGTCAGCTGCTGGCCCGTTC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6.7%; Score 30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
Disclosure, Col 18; 29pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP
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/*tag=
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hes 30; Conserv
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modified_base
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The present sequence is that of an oligonuclectide prepared using phosphoramidite chemistry. A nanoparticle conjugate was prepared by CC joining the 5'-meraploalkyl oligonuclectide to gold nanoparticles. The nanoparticle conjugate was used to enhance the sandwich amplification of a silver signal on glass slides containing silver sports from coligonucleotide assays performed using silver staining. This illustrated the method of the invention, which is based on the discoveries that: (1) gold nanoparticles coated with oligonucleotides bind to silver that has previously been deposited on gold nanoparticle-oligonucleotide conjugates commobilised by hybridisation on a glass substrate or plate; and (2) that the gold-nanoparticle-oligonucleotide-silver-gold oligonucleotide conjugates consorting silver deposition in detecting systems where the formation of silver spot serves as a reporter for the presence of amolecule, or including proteins, nucleic acids and small molecules. The detecting systems include detection of molecules in situ (e.g. on cells or in a tissue sample) and assays where the target molecules. The method has special cullity in increasing the signal strength in diagnostic and screening applications involving detection of target molecules arrayed at discrete companion in signal proteins in solid surface. It provides a means for greatly enhancing the sample, economical, and provides a large enhancement in signal
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6.7%; Score 30; DB 1; Length 30;
100.0%; Pred. No. 32;
ive 0; Mismatches 0; Indels
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97US-00833377.
99US-00420056.
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ABX10985 standard; DNA; 30
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Best Local Similarity 100.
Matches 30; Conservative
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04-APR-1997;
18-OCT-1999;
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This invention relates to a purified human telomerase protein, which when associated with telomerase RNA component has DNA polymerase activity.

Also disclosed in the specification is a method for assessing a regulator (preferably a telomerase inhibitor or activator of telomerase involves casuring telomerase enzymatic activity of the composition in presence of a regulator. The telomerase protein of the invention may be used in the activation of the invention may be used in the activity assays for measuring telomerase activity which are useful in characterising cancer and pre-cancer cells, for identifying and testing regulators of telomerase activity in in vitro assay and for preparing antibodies against telomerase. The mammalian telomerase protein of the invention is at least approximately 3000 fold more pure (in terms of telomerase activity per weight of protein) than a crude extract of telomerase activity per weight of protein) than a crude extract of cell from adenovirus- transformed kidney cell. Purified telomerase for facilitates a thorough biochemical analysis of the enzyme's mechanism for develophing mechanism- based regulators. The present sequence represents a human telomerase antisense oligonucleotide which has affinity to telomerase and is used to purify the telomerase protein of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pruzan RA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human telomerase antisense oligonucleotide primer oligol4.
                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1; Length 30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 30 BP; 6 A; 6 C; 8 G; 10 T; 0 U; 0 Other;
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telomerase protein with telomerase RNA component
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Pred. No. 32;
tive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 30;
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                                       Col 9; 24pp; English.
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97US-00833377.
99US-00420056.
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
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modified_base
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04-APR-1997;
18-OCT-1999;
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                                       Disclosure;
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ABX10986/c
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Also disclosed with telomerase RNA component has DNA polymerase activity.

Also disclosed in the specification is a method for assessing a regulator (preferably a telomerase RNA component has DNA polymerase activity.

Also disclosed in the specification is a method for assessing a regulator of preferably a telomerase inhibitor or activator of telomerase involves measuring telomerase involves measuring telomerase sorivity of the composition in presence of a regulator. The telomerase protein of the invention may be used in testing regulators of telomerase protein of the invention and carterising cancer and pre-cancer cells, for identifying and testing regulators of telomerase activity in in vitro assay and for preparing antibodies against telomerase. The mammalian telomerase protein of the invention is at least approximately 3000 fold more pure (in terms of telomerase activity per weight of protein) than a crude extract of cell from adenovirus- transformed kidney cell. Purified telomerase for facilitates a thorough biochemical analysis of the enzyme's mechanism for developing mechanism- based regulators. The present sequence represents a human telomerase antisense oligonucleotide which has affinity to
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invention relates to a purified human telomerase protein, which when
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention describes a method of obtaining mammalian telomerase protein (I). The method involves preparing enriched solution (ES) from a cell expressing telomerase where the component of (I) in ES is separated from other proteins expressed by cell by combining ES with oligonucleotide (O) having specific affinity for (I), and collecting
                                                                                                                                                                                                                                                                                                                                    telomerase and is used to purify the telomerase protein of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gape
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mammalian telomerase protein; telomerase purification; telomere; anion exchange matrix; selectivity matrix; gel filtration chromatography; gradient centrifugation; antisense oligonuclecide; se.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human telomerase RNA component antisense oligonucleotide seg id 5.
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                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Preu. ...
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97US-00833377.
99US-00420056.
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ADC35652/
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                                                                       containing telomerase protein with an anion exchange matrix, and collecting protein that binds the matrix, combining a fraction containing telomerase protein with a mation exchange matrix, and collecting protein that binds the matrix. Guch as a hegarin matrix), and collecting protein that binds the matrix. The method comprises successively enriching fractions containing telomerase protein on exchange matrices and combining a fraction containing telomerase protein with an intermediate selectivity matrix, collecting protein that binds the matrix, where the intermediate selectivity matrix and separating a fraction containing the telomerase protein by gel filtration chromatography or gradient centrifugation. The telomerase is enriched from an extract of cells stably expressing telomerase. This sequence represents an antisense oligonucleotide to the RNA component of human telomerase that can be used in the purification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Obtaining telomerase, by preparing enriched solution from cell expressing telomerase, combining the solution with oligonucleotide having specific affinity for the protein and collecting protein bound to oligonucleotide.
protein bound to (0). The oligonucleotide comprises a retrievable label such as biotin and contains a sequence that is specifically recognized by telomerase protein. The oligonucleotide contains or does not contain the sequence (TTAGGG)3. The method further comprises combining a fraction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention describes a method of obtaining mammalian telomerase protein (I). The method involves preparing enriched solution (ES) from a cell expressing telomerase where the component of (I) in ES is separated from other proteins expressed by cell by combining ES with
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        mammalian telomerase protein; telomerase purification; telomere; anion exchange matrix; selectivity matrix; gel filtration chromatography; gradient centrifugation; antisense oligonucleotide; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                          1; Length 30;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein (I). The method involves cell expressing telomerase where from other proteins expressed by
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97US-00833377.
99US-00420056.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADC35651 standard; DNA; 30
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                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity 100.
                                                                                                                                                                                                                                                                                                                                method of the invention.
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04-APR-1997;
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Matches
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RESULT 39
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          protein bound to (0). The oligonaclectide comprises a retrievable label such as biotin and contains a sequence that is specifically recognized by telemerase protein. The oligonaclectide contains or does not contain the sequence (TTMGGG)3. The method further comprises combining a fraction containing telomerase protein with an anion exchange matrix, and collecting protein that binds the matrix, combining a fraction containing approach with a cation exchange matrix. The method comprises successively enriching fractions containing telomerase protein that binds the matrix. The method of matrix), and collecting protein that binds the matrix. The method comprises successively enriching fractions containing telomerase protein that binds the matrix. The method on several different ion exchange matrices and combining a fraction containing telomerase protein with an intermediate selectivity matrix, collecting protein that binds the matrix, where the intermediate selectivity matrix, and separating a fraction containing the telomerase protein where the intermediate is selectively matrix and separating a fraction containing the telomerase protein by gel filtration chromatography or gradient centrifugation. The collecting protein the matrix of selectivity matrix and separating a fraction containing the telomerase is a function of the seal of t
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                                                                                                                                                                                                                                                                                                                            telomerase. This sequence represents an antisense oligonucleotide to the RNA component of human telomerase that can be used in the purification
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  oligonucleotide (0) having specific affinity for (I), and collecting
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100.0%; Pred. No. 5.,
... 0; Mismatches
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/mod_base= OTHER
/note= "Biotin labelled"
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04-APR-1997; 97US-00833377.
18-OCTT-1999; 99US-00420056.
20-NOV-2000; 2000US-00717828
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ATKINSON E M.
LICHTSTEINER S
VASSEROT A P.
PRUZAN R A.
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Best Local Similarity
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(LICH/)
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(PRUZ/)
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Identifying telomerase regulators useful for treating cancer.

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telomerase activity that may be useful for treating cancers. The method activity which may be useful for treating cancers. The method activity which may be useful as cancer treatments. It has been found that found that the cells of many human cancers have telomerase activity. This helps explain why cancer cells continue dividing without becoming senescent. If telomerase activity in cancer cells can be inhibited, the cancer cells are expected to reach senescence and cease dividing. The present sequence is human telomerase antisense oligonulceotide used to illustrate the method of the invention.
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                                                                                                                                                                                                                                                                                     Gaps
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                             invention relates to a method for identifying regulators of
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                                                                                                                                                                                                                                                 6.7%; Score 30; DB 1; Length 30; 100.0%; Pred. No. 32;
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Disclosure; SEQ ID NO 4; 22pp; English.
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/mod base= OTHER
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04-APR-1997; 97US-0083377.
18-OCT-1999; 99US-00420056.
20-NOV-2000; 2000US-0071828.
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ADG62874 standard; DNA; 30
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                                                                                                                                                                                                                                                                                    Conservative
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ATKINSON E M.
LICHTSTEINER S P
VASSEROT A P.
PRUZAN R A.
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es 30; Conserv
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(LICH/)
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activity which may be useful as cancer treatments. It has been found that found that the cells of many human cancers have telomerase activity. This helps explain why cancer cells continue dividing without becoming senescent. If telomerase activity in cancer cells can be inhibited, the cancer cells are expected to reach senescence and cease dividing. The present sequence is human telomerase antisense oligonulceotide used to illustrate the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; telomerase RNA component; anticancer therapy; purification; assay; vaccine; cancer; antisense oligonucleotide; ss.
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diagnosis and treatment of cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antisense oligonucleotide 5 for human telomerase RNA component.
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                                                                                                                                                                                    Query Match 6.7%; Score 30; DB 1; Length 30; Best Local Similarity 100.0%; Pred. No. 32; Matches 30; Conservative 0; Mismatches 0; Indels
                                                                                                                                                  Sequence 30 BP; 8 A; 5 C; 13 G; 4 T; 0 U; 0 Other;
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/note= "biotinylated"
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AAV63647 standard; DNA; 30 BP.
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modified_base
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                                                                                                                                                                                                                                                                                                                   human; immune response; cancer; vaccine; treatment; disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Compositions comprising human telomerase, useful for treating diseases
                                                                               Gape
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                                                   DB 1; Length 30;
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                                                                             Indele
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 associated with overexpression of telomerase e.g. cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pruzan
                                                                                                                                                                                                                                                                                       Human clone 28-1 telomerase oligonucleotide oligo-5.
                                                                                                                                                                                                                                                                                                                                                                                                                                  /*tag= a
/note= "5'-biotinylated guanosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 30 BP; 4 A; 11 C; 9 G; 6 T; 0 U; 0 Other;
                        Seguence 30 BP; 4 A; 11 C; 9 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Weinrich SL,
                                                 Score 28.4; DB Pred. No. 45; 0; Mismatches
                                                                                                         GAGCTGTGGGACGTGCACCCAGGACTCGGC 441
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GAGCTGTGGGACGTGCACCCCAGGACTCGGC 441
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6.3%; Score 28.4;
16.7%; Pred. No. 45;
                                                                                                                                 GAGCTATGGGACGTGCACCCAGGACTCGGC 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Col 43-44; 34pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lichtsteiner SP,
                                                  6.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              97US-00833377.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          95US-00510736
                                                                                                                                                                                                       AAZ23629 standard; DNA; 30
                                                                                                                                                                                                                                                            (first entry)
                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                /*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  duery Match
Best Local Similarity
Matches 29; Conserv
                                            Query Match
Best Local Similarity
Matches 29; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GERO-) GERON
                                                                                                                                                                                                                                                                                                                                                                                                                 nodified base
                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          04-AUG-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ΕΜ,
AP;
                                                                                                                                                                                                                                                             07-JAN-2000
                                                                                                                                                                                                                                                                                                                   Telomerase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US5968506-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-OCT-1999
                                                                                                                                                                                                                                                                                                                                   primer; ss.
                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                       412
                                                                                                                                                                                                                                  AAZ23629;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Vasserot
cancer
                                                                                                                                                                              RESULT 41
                                                                                                                                                                                          AAZ23629,
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US09436060A-16.rng.sl

Human telomerase antisense oligonucleotide primer oligos.

(first entry)

17-AUG-2003

ABX10984;

ABX10984 standard; DNA; 30 BP.

RESULT 43 ABX10984/c

relomerase; antisense; primer; oligo5; ss; cancer.

/*tag= a /note= "Biotinylated" Location/Qualifiers

Key modified_base

Synthetic

US6517834-B1

11-FEB-2003

20-NOV-2000; 2000US-00717828.

04-AUG-1995;

04-APR-1997;

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The sequence represents a biotinylated antisense oligonucleotide used in the purification of human telomerase. The invention relates to a purified human telomerase core enzyme protein comprising 2000-fold increased purity compared with a crude extract of cells from adenovirus-transformed kidney cell line (293 cells) and when associated with telomerase RNA component has DNA polymerase activity and a molecular weight of 200-2000 kilo Daltons (kDa). The purified telomerase is useful for inducing a humoral or cell-mediated immune response in an animal. Purified telomerase or immunogenic fragments are useful as vaccines for treating diseases associated with over-expression of telomerase, such as cancer and for producing antibodies that recognize telomerase, which are useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   as affinity agents in isolating the proteins and for detecting the presence of proteins in a sample, such as cell or tissue. Identification of telomerase aids in diagnosis of cancer or pre-cancerous states. Telomerase and/or telomerase associated proteins are also useful for screening compounds to identify agents that alter the association of telomerase-associated proteins, such as nucleolin or EF2H with telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Purified human telomerase, useful for inducing immune response in animals, comprises several thousand folds increased purity compared with cytoplasmic crude cell preparations.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Atkinson EM, Lichtsteiner SP, Vasserot AP, Pruzan RA;
                                                                                                                                                                                                                                                                                                                                                  Human, Telomerase, vaccine, antibody, cancer, BF2H, nucleolin, antisense oligonucleotide, Oligo 5, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6.3%; Score 28.4; DB 1; Length 30; 96.7%; Pred. No. 45; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                   Antisense oligonucleotide for human telomerase, Oligo 5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seguence 30 BP; 4 A; 11 C; 9 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /*tag= a
/mod_base= G
/note= "G is biotinylated"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
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                                                                  AAS09474 standard; DNA; 30 BP.
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97US-00833377
                                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Key
modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
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04-APR-1997;
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                                                                                                                                                                                                            24-OCT-2001
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                                                                                                                                     AAS09474;
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                          AASO9474/C
XX
XX
AACO9474/C
XX
AACO
XXX
RESULT 42
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associated with telomerase RNA component has DNA polymerase activity.

Also disclosed in the specification is a method for assessing a regulator (prefeated) at elomerase inhibitor or activator of telomerase involves measuring telomerase enzymatic activity of the composition in presence of a regulator. The telomerase protein of the invention may be used in developing and testing assays for measuring telomerase activity which are useful in characterising cancer and pre-cancer cells, for identifying and testing regulators of telomerase activity in in vitro assay and for preparing antibodies against telomerase. The mammalian telomerase protein of telomerase activity per weight of protein) than a crude extract of cell from adenovirus- transformed kidney cell. Purified telomerase (a facilitates a thorough biochemical analysis of the enzyme's mechanism for developing mechanism- based regulators. The present sequence represents a thomorase autisense oligonuclocide which has affinity to thomorase and is used to purify the telomerase protein of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ô
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention relates to a purified human telomerase protein, which when
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pruzan RA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Composition useful e.g. in diagnosis of cancer comprises complex of telomerase protein with telomerase RNA component.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .,
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Weinrich SL, Atkinson EM, Lichtsteiner SP, Vasserot AP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6.3%; Score 28.4; DB 1; Length 30; 96.7%; Pred. No. 45;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 30 BP; 4 A; 11 C; 9 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 13; Col 9; 24pp; English.
                                                                                                                                                                                                                                                                                                                                                      95US-00510736.
97US-00833377.
99US-00420056.
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                                                                                                                                                                                                                                                                                                                                                                                                                         (GERO-) GERON CORP.
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Matches 29; Conserv
                                                                                                                                                                                                                                                                                                                                                                                         18-OCT-1999;
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ID ADC356
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Gaps ; 0

412 GAGCTGTGGGACGTGCACCCAGGACTCGGC 441

Local Similarity 96.7 nes 29; Conservative

Matches

à 엄

GAGCTATGGGACGTGCACCCAGGACTCGGC 1

30

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Telomerase activity; therapy; cancer; cytostatic; antisense; ss.
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                                                                                                                                 Human telomerase RNA antisense oligonucleotide, oligo 5.
  ADG62872 standard; DNA; 30 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      04-APR-1997; 97US-00833377
18-OCT-1999; 99US-00420056
20-NOV-2000; 2000US-00717828
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24-DEC-2002; 2002US-00330872
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Weinrich SL, Atkinson EM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WEINRICH S L.
ATKINSON E M.
LICHTSTEINER S P
VASSEROT A P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-811733/76.
                                                                                                                                                                                                                                                                                                                                                                                                    US2003186282-A1
                                                                                                                                                                                                                                                                                       modified base
                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-AUG-1995;
04-APR-1997;
                                                                                     11-MAR-2004
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(VASS/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (PRUZ/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (WEIN/)
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention describes a method of obtaining mammalian telomerase protein (1). The method involves preparing enriched solution (ES) from a cell expressing telomerase where the component of (1) in ES is separated from other proteins expressed by cell by combining ES with oligonucleotide (0) having specific affinity for (1), and collecting protein bound to (0). The Oligonucleotide comprises a retrievable label such as biotin and contains a sequence that is specifically recognized by telomerase protein. The Oligonucleotide contains or does not contain the sequence (TTAGGG)3. The method further comprises combining a fraction containing telomerase protein with a cation exchange matrix, and collecting protein with a cation exchange matrix. The method collecting protein with a cation exchange matrix. The method containing a fraction containing telomerase protein with a cation exchange matrix. The method comprises successively enriching fractions containing telomerase protein with that binds the matrix. The method containing telomerase protein with an intermediate selectivity matrix, collecting protein that binds the matrix, where the intermediate selectivity matrix, collecting protein that binds the matrix, where the intermediate protein and separating a fraction containing the centrification. The elomerase in enriched from an extract of cells stably expressing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Obtaining telomerase, by preparing enriched solution from cell expressing telomerase, combining the solution with oligonucleotide having specific affinity for the protein and collecting protein bound to oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         telomerase. This sequence represents an antisense oligonucleotide to the RNA component of human telomerase that can be used in the purification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Weinrich SL, Atkinson EM, Lichtsteiner SP, Vasserot AP, Pruzan RA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                    mammalian telomerase protein; telomerase purification; telomere; anton exchange matrix; selectivity matrix; gal filtration chromatography; gradient centrifugation; antisense oligonucleotide; ss.
                                                                                                          Human telomerase RNA component antisense oligonucleotide seq id
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           6.3%; Score 28.4; D 96.7%; Pred. No. 45; ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GAGCTATGGGACGTGCACCCAGGACTCGGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 4; SEQ ID NO 3; 24pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                           95US-00510736.
97US-00833377.
99US-00420056.
                                                                                                                                                                                                                                                                                                                                                                                                  20-NOV-2000; 2000US-00717829
                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GERO-) GERON CORP.
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Best Local Similarity
Matches 29; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                           04-AUG-1995;
04-APR-1997;
18-OCT-1999;
                                                                                                                                                                                                                                                                                                            US6545133-B1
                                                                18-DEC-2003
                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                      08-APR-2003.
                       ADC35650;
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Pruzan RA;

Vasserot AP,

Lichtsteiner SP,

/*tag= a /mod_base= OTHER /note= "Biotin labelled"

95US-00510736

Location/Qualifiers

(first entry)

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                                           The invention relates to a method for identifying regulators of telomerase activity that may be useful for treating cancers. The method may be useful for treating cancers. The method may be useful as cancer treatments of telomerase activity which may be useful as cancer treatments. It has been found that found that the cells of many human cancers have telomerase activity. This sensescent. If telomerase activity in cancer cells can be inhibited, the cancer cells are expected to reach sensescence and cease dividing. The present sequence is human telomerase antisense oligonulceotide used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gарв
                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                            DB 1; Length 30;
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                                                                                                                                                                                                                                                                                                                                          Sequence 30 BP; 4 A; 11 C; 9 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      412 GAGCTGTGGGACGTGCACCCAGGACTCGGC 441
                                                                                                                                                                                                                                                                                                                                                                                                                 Pred. No. 45;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GAGCTATGGGACGTGCACCCAGGACTCGGC 1
                                                                                                                                                                                                                                                                                                                                                                                            6.3%; Score 28.4;
                                                                                                                                                                                                                                                                                      illustrate the method of the invention.
Claim 11; SEQ ID NO 3; 22pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                 ilarity 96.7%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAX90787 standard; DNA; 28
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les 29, Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAX90787;
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ID AAX9
XX
AC AAX9
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RESULT 45 ADG62872/c

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03-AUG-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAX77130;
       Kim NW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Funk W;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAX77130/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 48
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (hTR). It is used to amplify human staufen (hStau) cDNA synthesised using random hexamer primers and Superscript II reverse transcriptase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RNA component; human telomerase; antisense oligonucleotide; infection; neuroblastoma; bladder cancer; colon cancer; prostate cancer; cancer; contraception; sterilisation; immunosuppression; therapeutic; hTR; immune system down-regulation; anti-inflammatory therapy; RT-PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Human staufen polypeptide useful in methods for identifying telomerase inhibitors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present sequence is a PCR primer specific to human telomerase RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                      PCR primer; human telomerase RNA; hTR; amplify; human staufen cDNA; hStau; synthesised; random hexamer primer; Superscript II reverse transcriptase; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RNA component of human telomerase (hTR) amplifying reverse primer.
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Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 28 BP; 2 A; 4 C; 13 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Prea. ...
                                                                                                                                                                                                                                                                                                                    (UYJO ) UNIV JOHNS HOPKINS SCHOOL MEDICINE.
Human telomerase RNA specific PCR primer-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17 GCCTGGGAGGGGTGGTGGCCATTTTTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 15; 50pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAV41193 standard; DNA; 27 BP
                                                                                                                                                                                                                                              99WO-US007533
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96US-00770565
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                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-620168/53
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                                                                                                                                                                                                                                                                                                                                                           Le S;
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                                                                                                                                  Homo sapiens.
                                                                                                                                                                       WO9951255-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9828442-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                           Greider CW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       08-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-DEC-1996;
                                                                                                                                                                                                          14-0CT-1999
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This primer is used for the RT-PCR amplification of an RNA component of human telomerase (hTR). This is used in the method of invention of determining the amount of hTR in a sample. The method comprises camplifying a sequence of hTR and a control polynucleotide from a sample and determining an amount of amplified hTR and an amount of amplified control polynucleotide. The amount of amplified hTR is normalised with crespect to the amount of Amplified control polynucleotide to provide a normalised amount of hTR which provides antisense oligonucleotides to the hTR which may specifically be used for detection of an RNA component of human telomerase in a sample. This is useful for diagnosing cancer (especially neuroblastoma, bladder, colon and prostate cancer), and providing prognosis for a cancer patient. The antisense oligonucleotides con bused for inhibiting telomerase activity in both cultured cells and in cells in vivo. They can be used in therapeutics for treating or contraception or sterilisation, for immunosuppression, and for selectively down-regulating specific branches confinammatory theraples or for treating infections by, e.g. yeast,
                                                                                                                                    New polynucleotide(s) anti:sense to human telomerase - used for detecting or inhibiting human telomerase, e.g. for treating cancers, contraception, immuno-suppression or treating infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New human GC6 gene, useful for identifying agents for treating diseases and/or conditions associated with cell senescence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cellular senescence; modulator; GC6 gene; senescent gene expression;
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00.0%; Pred. No. 51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
Pruzan R, Weinrich SL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 27 BP; 7 A; 3 C; 10 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; Pred. ....
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  144 CCTTCCACGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27 cerrecacerrearreragadeaaac 1
                                                                                                                                                                                                                                                                                               Claim 65; Page 75; 80pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     98WO-US024996.
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Wu F, Kealey JT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAX77130 standard; DNA; 27
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Best Local Similarity 100.4
Matches 27, Conservative
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                                                                      WPI; 1998-377670/32.
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Length 27;

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                                                               senescence. Recombinant expression vectors comprising a recombinant polynucleotide corresponding to a polynucleotide in a human GCG gene, are useful for altering senescent gene expression. The vectors and host cells comprising the protein gene expression. The polynucleotides are useful for producing the protein, pGCG and nucleic acid derivatives. The proteins encoded are useful for rational antibodies specific for pGCG, which are useful for isolating pGCG, and for detecting cells comprising pGCG in complex cell mixtures. The characterization of the polynucleotides enable the identification of therapeutic agents that identify and distinguish between young and senescent cells. This enables treatment of aging diseases induced or exacerbated by cellular senescence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to a method for diagnosing and/or monitoring the evolution of cancers. The method comprises analysing enzyme telomerase RNA in blood plasma or serum. The method is typically used for diagnosing breast, ovarian, stomach or colon cancer and/or monitoring the evolution of the cancers after treatment by chemotherapy or operations. The present sequence is a PCR primer for human telomerase RNA (hTR), which was used in the example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    e.g. breast
plasma or serum.
                                                  The invention relates to methods for modulating and identifying cellular
                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; telomerase RNA; PCR primer; cancer; breast; ovarian; stomach;
colon; hTR; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       e, reliable diagnosis and/or prognosis of cancer, by analyzing the RNA components of telomerase in
                                                                                                                                                                                                                                                                                              Sequence 27 BP; 4 A; 6 C; 11 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human telomerase RNA, hTR, antisense PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                     425 TGCACCCAGGACTCGGCTCACACATGC 451
                                                                                                                                                                                                                                                                                                                                                                                                                         27 TGCACCCAGACTCGGCTCACACATGC 1
                  5; Page 74; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-MAY-2000; 2000EP-00111370.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26-MAY-2000; 2000EP-00111370
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ABA95497 standard; DNA; 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (CHEN/) CHEN X Q. (STRO/) STROUN M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ANKER P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12-MAR-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABA95497;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (STRO/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 49
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Sequence 27 BP; 7 A; 3 C; 10 G; 7 T; 0 U; 0 Other;

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The invention relates to promoter regions from mouse and human telomerase

RNA (TR) component genes. The TR gene promoter can be linked to a

RNA (TR) component genes. The TR gene encoding a cytocoxin, for therapy of

cancer, especially neoplasia. The telomerase is necessary for the

cancer, especially neoplasia. The telomerase is necessary for the

currestricted proliferative capacity of many human cancers. Mutation or

dysregulation of the telomerase expression pathway may cause reactivation

or upregulation of telomerase expression in cancer. Substances,

cleantified in the methods, can be used to block transcription from the TR

gene promoter through interaction of the 5' regulatory sequences. These

cuspicide nucleic acids and factors that disrupt signal transduction, are

cuseful for cancer therapy. In particular, gene therapy vectors

cuseful for cancer therapy. In particular, gene therapy vectors

cuseful for cancer therapy. In particular, gene therapy vectors

cuseful for cancer therapy. In particular, gene therapy vectors

cuseful for cancer therapy in particular, gene therapy vectors

cuseful for cancer therapy in particular, gene therapy vectors

cuseful for cancer therapy and the promoter and a viral thymidine

cuseful for an be controlled or treated. Direct down-regulation of

ceffective anticancer therapy and the cloning of the hTR gene promoter

allows the analysis of therapeutic molecules which modulate hTR promoter

callows the analysis of therapeutic molecules which modulate hTR promoter

callows the analysis of therapeutic molecules which modulate for amplifying
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                                 Gapв
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gape
                                                                                                                                                                                                                                                                                                                                                                                                                      Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; gene therapy; thymidine kinase gene; anticancer therapy; human;
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                                               0; Indela
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                           Human telomerase RNA gene (hTR) specific primer hTR14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 30 BP; 5 A; 8 C; 8 G; 9 T; 0 U; 0 Other;
    6.0%; Score 27; DB
ilarity 100.0%; Pred. No. 51;
Conservative 0; Mismatches
                                                                                          144 CCTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CANC-) CANCER RES CAMPAIGN TECHNOLOGY.
                                                                                                                         CCTTCCACCGTTCATTCTAGAGCAAAC 1
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                                                                                                                                                                                                                             264/c
AAZ07264 standard; DNA; 30
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Query Match
Best Local Similarity
Matches 27; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               human TR gene (hTR)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
Matches 28, Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-JAN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                          AAZ07264;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Keith WN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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mammal; antisense; triplex forming oligonucleotide; plasmid;

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The RNA components of (non) human mammalian telomerase(s) especially from mouse, rat and chinese hamster are all claimed. Antisense oligonucleotides can be used to block the activity of the telomerase; probes and primers can be used in detection; vectors and host cells transformed with the isolated telomerase genes can be used for production of telomerases; RNA and DNA ribozymes and triplex forming oligonucleotides directed against the telomerase agnes can be used therapeutically as can plasmids. A mouse which lacks the telomerase gene (also claimed) can be used for study of telomere regulation in vivo, and hard role it plays in immortalisation. Three primers (AAT11040, AAT11043, AAT11044) were used to produce antisense oligonucleotides which were then used to produce antisense oligonucleotides which were then alongside both AAT11043 and AAT11044 to produce two different antisense
                                                                   Primer for production of telomerase antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      senescence and immortalisation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 8; Page 53; 85pp; English.
                                                                                                                  probe; primer; ribozyme; ss.
                                     02-JUL-1996 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1996-097428/10.
                                                                                                                                                                                                                                                                                                                                                              GERON CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RNA components of
                                                                                                                                                                            WO9601614-A2
                                                                                                                                                                                                                                      07-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                COLD
                                                                                                                                                                                                                                                                                                                                                                                                          Villeponteau
                                                                                                                                                                                                                                                                    07-JUL-1994;
                                                                                                                                                                                                                                                                                                                 07-JUN-1995;
                                                                                                  Telomerase;
                                                                                                                                                                                                                                                                                    27-OCT-1994;
13-FEB-1995;
                                                                                                                                                                                                         25-JAN-1996
                                                                                                                                                                                                                                                                                                                                                                                           Andrews WH,
                                                                                                                                             Synthetic.
         AAT11044;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                              (COLD-)
                                                                                                                                                                                                                                                                                                                                                              (GERO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cell
        à
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence, a forward primer for the RNA component of human telomerase (RCHT), was used in a primed in situ labelling (PRINS) procedure. The RCHT can be used in the recombinant prodn. of an active telomerase mol., capable of adding sequences to chromosomal DNA telomeras, and in the synthesis of mutant sequences for the detection of mutant mammalian telomerase RNA component polymoleotides. The RCHT may also be used in the identification of telomerase modulating agents, and in the detection of telomerase related, or neoplastic conditions in a patient. Polymoleotides of at least 25 consecutive mucleotides identical, or complementary to the RCHT sequence linked to heterologous transcriptional regulatory sequences, can be used for the gene therapy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RNA component of mammalian telomerase, esp. human - useful in identifying
                                                                                                                                                                                                                          RNA component; human; telomerase; forward primer; PRINS; recombinant production; synthesis; mutant; detection; mammalian; identification; modulating agent; neoplastic condition; transcriptional regulatory sequence; gene therapy; disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1; Length 28;
                                                                                                                                                                                               RNA component of human telomerase PRINS forward primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 28 BP; 2 A; 3 C; 13 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Andrews WH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     e.g. candidate telomerase-modulating agents.
                   CTAACCCTAACTGAGAGGGCGTAGGATCC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5.9%; Score 26.4; 96.4%; Pred. No. 60
CTAACCCTAACTGAGAAGGGCGTAGGCGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 13; Page 91; 114pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Funk W,
                                                                                                          BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            94US-00330123.
95US-00472802.
95US-00482115.
                                                                                                                                                                                                                                                                                           primed in situ labelling; ss
                                                                                                                                                                                                                                                                                                                                                                                                                    95WO-US008530
                                                                                                                                                                                                                                                                                                                                                                                                                                                94US-00272102
                                                                                                       AAT10309 standard; DNA; 28
                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Feng J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1996-097581/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
Les 27; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Villeponteau B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   human diseases
                                                                                                                                                                                                                                                                                                                                                      WO9601835-A1
                                                                                                                                                                                                                                                                                                                                                                                                                  06-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                             07-JUL-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-OCT-1994;
07-JUN-1995;
                                                                                                                                                                    10-SEP-1996
                                                                                                                                                                                                                                                                                                                                                                                   25-JAN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-JUN-1995
                                                                                                                                                                                                                                                                                                                         Synthetic.
                           30
                                                                                                                                     AAT10309;
 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                        RESULT 51
                                                                                                                     à
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(non)human mammalian telomerase(s) - useful in studying

Marhuenda MA;

Greider C,

Funk W,

Feng J,

Avillon AA,

94US-00272102. 94US-00330123. 95US-00387524. 95US-00485778.

95WO-US008620

SPRING HARBOR LAB

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                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                   RNA component; human; telomerase; polymerase chain reaction; recombinant production; synthesis; mutant; detection; mammalian;
                                                                                      ó
                                              score 26; DB 1; Length 26;
Pred. No. 59;
                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                      RNA component of human telomerase nested PCR primer R3c.
                        Sequence 26 BP; 7 A; 3 C; 9 G; 7 T; 0 U; 0 Other;
                                            5.8%; Scc. 100.0%; Pred. No. ...
                                                                                                                145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                            26 criccaccerrearicragaccaaac 1
                                                                                                                                                                                                                      BP
                                                                                                                                                                                                      304/c
AAT10304 standard; DNA; 26
                                                                                                                                                                                                                                                                             (first entry)
                                                                     Similarity 100.
26; Conservative
                                                                                                                                                                                                                                                                            10-SEP-1996
molecules
                                                                                                                                                                                                                                                AAT10304;
                                                                                                                                                                                                   AAT10304/c
ID AAT10
XX
AC AAT10
XX
XX
DE RNA C
XX
XX
KW RNA C
KW RWA C
                                                                                                                                                                                         RESULT 53
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0

Gaps

. 0

1; Indels

0; Mismatches

Conservative

Matches

44 28

GCCTGGGAGGGGTGGTGGCTATTTTTG 17 GCCTGGGAGGGGTGGTGGCCATTTTTG

ద ò

AAT11044 standard; DNA; 26 BP.

AAT11044/c ID AAT1104 XX

RESULT 52

95WO-US008530.

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(GERO-) GERON
                                      07-JUL-1994;
27-OCT-1994;
07-JUN-1995;
07-JUN-1995;
  06-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9601835-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                06-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27-OCT-1994;
07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-JUL-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-JAN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAT10306;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAT10306
  g
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence, a nested PCR primer for the RNA component of human telomerase (RCHT), was used in a 5' RACE procedure. The RCHT can be used in the recombinant prodn. of an active telomerase mol., capable of adding sequences to chromosomal DNA telomeras, and in the synthesis of mutant sequences for the detection of mutant mammalian telomerase RNA component polynucleotides. The RCHT may also be used in the identification of telomerase modulating agents, and in the detection of telomerase related, or neoplastic conditions a patient. Polynucleotides of at least 25 consecutive nucleotides identical, or complementary to the RCHT sequence linked to heterologous transcriptional regulatory sequences, can be used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RNA component of mammalian telomerase, esp. human - useful in identifying
modulating agent; neoplastic condition;
regulatory sequence; gene therapy; disease; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RNA component; human; telomerase; lung fibroblast; cell line Wi 38; recombinant production; synthesis; mutant; detection; mammalian; dentification; mandulating agent; neoplastic condition; transcriptional regulatory sequence; gene therapy; disease; polymerase chain reaction; antisense plasmid; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RNA component of human telomerase antisense plasmid PCR primer R3C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5.8%; Score 26; DB 1; Length 26;
100.0%; Pred. No. 59;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 26 BP; 7 A; 3 C; 9 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                            Funk W, Andrews WH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                e.g. candidate telomerase-modulating agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Preα. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gene therapy of human diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CTTCCACCGTTCATTCTAGAGCAAAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 10; Page 82; 114pp; English
                                                                                                                                                                                                                                                                94US-00330123.
95US-00472802.
95US-00482115.
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                                                                                                                                                                                                                                               94US-00272102
                                                                                                                                                                                                                                                                                                                                                                                          Villeponteau B, Feng J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1996-097581/10.
                                                                                                                                                                                                                                                                                                                                                   (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
les 26; Conserv
identification;
transcriptional
                                                                                                                       WO9601835-A1
                                                                                                                                                                                                      06-JUL-1995;
                                                                                                                                                                                                                                                                                                           07-JUN-1995;
                                                                                                                                                                                                                                             07-JUL-1994;
                                                                                                                                                                                                                                                                  27-OCT-1994;
07-JUN-1995;
                                                                                                                                                                25-JAN-1996
                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
KEYER 
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                                                                                                                                                                                                                                                                                                                                                                   RNA component of mammalian telomerase, esp. human - useful in identifying e.g. candidate telomerase-modulating agents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RNA component; human; telomerase; polymerase chain reaction; recombinant production; synthesis; mutant; detection; madulating agent; neoplastic condition; modulating agent; neoplastic condition; transcriptional regulatory sequence; gene therapy; disease; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gapв
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5.8%; Score 26; DB 1; Length 26;
100.0%; Pred. No. 59;
iive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 26 BP; 7 A; 3 C; 9 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RNA component of human telomerase PCR primer F3b.
                                                                                                                                                                                                                                     Andrews WH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      145 CTTCCACCGTTCATTCTAGAGCAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CTTCCACCGTTCATTCTAGAGCAAAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 8; Page 80; 114pp; English.
                                                                                                                                                                                                                                     Funk W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  94US-00272102.
94US-00330123.
95US-00472802.
95US-00482115.
94US-00272102.
94US-00330123.
95US-00472802.
95US-00482115.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       95WO-US008530.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAT10306 standard; DNA; 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 100.0
Matches 26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-SEP-1996 (first entry)
                                                                                                                                                                                                                                     Feng J,
                                                                                                                                                                                                                                                                                                           WPI; 1996-097581/10.
                                                                                                                                                                       CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (GERO-) GERON CORP.
                                                                                                                                                                                                                                     Villeponteau B,
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim
                                                                                                                                                                                                                                                                                                                                                                     RESULT 57
AAV41192
8X888888888888X8
                                                                                                                                                                                                                                                                                             ò
                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                     The present sequence, a PCR primer for the RNA component of human telomerase (RCHT), was used in a 3' RACE procedure. The RCHT can be used in the recombinant prodn. of an active telomerase mol., capable of adding sequences to chromosomal DNA telomeres, and in the synthesis of mutant sequences for the detection of mutant mammalian telomerase RNA component polynucleotides. The RCHT may also be used in the identification of telomerase modulating agents, and in the detection of telomerase related, or neoplastic conditions in a patient. Polynucleotides of at least 25 consecutive nucleotides identical, or complementary to the RCHT sequence linked to heterologous transcriptional regulatory sequences, can be used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the
                                                           esp. human - useful in identifying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cancer; eukaryotic parasite; hTR; vertebrate telomerase; yeast; protozoa; tumour; antibody; polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA encoding essential RNA components of human telomerase - also truncated or recombinant telomerase, useful for diagnosis and treatment of cancer and infection by eukaryotic parasites.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence represents PCR 3'-primer R3C used for amplifying human telomerase (hTR). The RNA and DNA can be used in hybridisation assays to detect or quantify telomerase activity in cells, tissue or fluid samples, e.g. for diagnosis of eukaryotic parasites (yeast and protozoa) or tumours. It is also useful as primers for amplification assays. The truncated or recombinant vertebrate telomerase is used
                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                     ;
0
                                                                                                                                                                                                                                                                                                                                                       5.8%; Score 26; DB 1; Length 26; 100.0%; Pred. No. 59;
                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
                                                                                                                                                                                                                                                                                                                           Sequence 26 BP; 8 A; 6 C; 7 G; 5 T; 0 U; 0 Other;
   Andrews WH;
                                                            RNA component of mammalian telomerase, esp. e.g. candidate telomerase-modulating agents.
                                                                                                                                                                                                                                                                                                                                                                  100.0%; Prec.
                                                                                                                                                                                                                                                                                             for the gene therapy of human diseases
                                                                                                                                                                                                                                                                                                                                                                                                                   45 TCTAACCCTAACTGAGAAGGGCGTAG 70
                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 10; Page 83; 114pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human telomerase PCR 3'-primer R3C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 5; Page 32; 48pp; English.
 Funk W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (COLD-) COLD SPRING HARBOR LAB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           96WO-US009517.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         95US-00478352.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAT58811 standard; DNA; 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
   Villeponteau B, Feng J,
                                                                                                                                                                                                                                                                                                                                                                                       26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Greider C, Autexier C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1997-099928/09.
                                WPI; 1996-097581/10
                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-JUN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9640868-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-NOV-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAT58811;
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screening assay) while the RNA or other inhibitors such as antisense molecules, are used to reduce such activity. Typical applications are initiation/restoration of activity to cause senescence or to prevent immortalisation of calls in tunours or parasites. The DNA is also used to produce recombinant telomerase, which can then be used conventionally to raise antibodies for diagnostic detection of telomerase. Detecting telomerase allows early diagnosis of tumour or infection, before clinical signs manifest. Telomerase inhibitors directed against e.g. Trypanosoma should cause fewer side effects than drugs currently used to treat such infections. The DNA encodes those parts of hTR RNA essential for activity but are significantly shorter than the endogenous RNA component
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RNA component; human telomerase; antisense oligonucleotide; infection; neuroblastoma; bladder cancer; colon cancer; prostate cancer; cancer; contraception; sterilisation; immunosuppression; therapeutic; hTR; immune system down-regulation; anti-inflammatory therapy; RT-PCR; primer;
therapeutically to increase telomerase activity (also as reagents in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This primer is used for the RT-PCR amplification of an RNA component of human telomerase (hTR). This is used in the method of invention of determining the amount of hTR in a sample. The method comprises amplifying a sequence of hTR and a control polynucleotide from a sample and determining an amount of amplified hTR and an amount of amplified are control polynucleotide. The amount of amplified of the sequence of hTR is normalised with respect to the amount of amplified control polynucleotide to provide anormalised amount of hTR which provides a determination of the amount of hTR in the sample. The invention provides antisense oligonucleotides to
                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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Les 26; Conservative
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the hTR which may specifically be used for detection of an RNA component of human telomerase in a sample. This is useful for diagnosing cancer (especially neuroblastoma, bladder, colon and prostate cancer), and providing prognosis for a cancer patient. The antisense oligonucleotides can be used for inhibiting telomerase activity in both cultured cells and in cells in vivo. They can be used in therapeutics for treating or preventing cancer, for contraception or sterilisation, for immunosuppression, and for selectively down-regulating specific branches of the immune system, e.g. a specific subset of T-cells, in anti-inflammatory therapies or for treating infections by, e.g. yeast,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; telomerase reverse transcriptase; hTRT; TRT; diagnosis; prognosis; cell proliferation; cancer; ageing; ribonucleoprotein; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence represents a PCR primer from the present invention which describes human telomerase reverse transcriptase (hTRT). The present invention also describes the following methods. (A) determining whether a test compound is a modulator of hTRT, by detecting the change in hTRT recombinant protein or polynucleotide, on administration of the
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970S-00854050.
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compound; (B) preparation of recombinant telomerase by contacting a protein preparation of HTRT with a telomerase RNA component; (C) detection of the HTRT RNA or protein in a sample by binding a relevant probe to the sample and detecting the complex formed or in the case of RNA detection, amplifying the product and correlating the presence of complex or amplification product with presence of HTRT in the sample; and (D) increasing the proliferation of a vertebrate cell by increasing hTRT expression; and (E) the use of an agent that causes an increase in cell vertebrate cell proliferation to create a medicament that inhibits ageing. A protein preparation of hTRT and the polynucleotide encoding hTRT can be used in the manufacture of medicaments for inhibiting the effect of ageing or cancer. Inhibitors of telomerase activity can be used to treat conditions that are associated with high telomerase activity. A protein preparation of hTRT can also be used in the new methods
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100.0%; Pred. No.
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97US-00846017.
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present invention also describes the following methods: (A) determining whether a test compound is a modulator of hTRT, by detecting the change in hTRT recombinant protein or polymucleotide, on administration of the compound; (B) preparation of recombinant telomerase by contacting a protein preparation of hTRT with a telomerase RNA component; (C) detection of the hTRT RNA or protein in a sample by binding a relevant probe to the sample and detecting the complex formed or in the case of RNA detection, amplifying the product and correlating the presence of complex or amplification product with presence of hTRT in the sample; and (D) increasing the proliferation of a vertebrate cell by increasing hTRT expression; and (E) the use of an agent that causes an increase in cell vertebrate cell proliferation of reate a medicament that inhibits of ageing A protein preparation of hTRT and the polymucleotide encoding hTRT can be used in the manufacture of medicaments for inhibiting the effect of ageing or cancer. Inhibitors of telomerase activity can be used
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                                                                                                                                                                                                                                                                                    treat conditions that are associated with high telomerase activity. A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene; TPC2; TPC3; telomere length; telomerase; human; cancer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAV19488;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   amplification of the human telomerase hTR gene (see AAV19481). hTR mRNA levels were showed to correlate with telomerase activity levels in a variety or mortal and immortal cell lines. Methods of the invention allow detection and quantitation of TPC2 (see AAV19479), TPC3 (see AAV19480) and/or TPC2 gene products and can be used to detect immortal cells, especially telomerase positive cancer cells
                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                   TPC3 and TR genes - regulate telomere length or modulate
                                                                                                                                                                                                                    hTR gene; TPC2; TPC3; telomere length; telomerase; human; cancer; gene therapy; diagnosis; PCR; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            and F3b (see AAV19488) were designed for the PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1; Length 26;
59;
Length 26;
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                      Indels
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                                                                                                                                                                                                                                                                                                                                                                                                         Adams RR;
 DB 1;
59;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human telomerase RNA specific PCR primer-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 26;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         145 CTTCCACCGTTCATTCTAGAGCAAAC 170
 Score 26;
Pred. No.
                                               70
                                                                    1 reraacceraacreagaagegegrag 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTTCCACCGTTCATTCTAGAGCAAAC 1
                                                                                                                                                                                                                                                                                                                                                                                                         Andrews WH,
                                               45 TCTAACCCTAACTGAGAAGGGCGTAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 49; 86pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5.8%; Sc.
100.0%; Pred
0; M
                                                                                                                                                                                                 Human hTR gene RT-PCR primer R3c.
 5.8%; Sccilarity 100.0%; Pr
Conservative 0;
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                                                                                                                             ВР
                                                                                                                                                                                                                                                                                                                                       96WO-US014679.
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AAX90788 standard; DNA; 26
                                                                                                                             AAV19489 standard; DNA; 26
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                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                         Feng J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  telomerase activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1998-207373/18
  Query Match
Best Local Similarity
Matches 26; Conserv
                                                                                                                                                                                                                                                                                                                                                                                   (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                         Villeponteau B,
                                                                                                                                                                                                                                                                    Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human TPC2,
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                                                                                                                                                                                                                                                                                                                 19-MAR-1998
                                                                                                                                                                                                                                                         Synthetic
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                                                                                                                                                    AAV19489;
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                                                                                                                     AAV19489/c
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                                                                                                       RESULT 61
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telomeric repeat amplification protocol.
                                                                                                                                                                                                                                                                                                                                                            PCR primer; 89.
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                                                                                                                                                                                                                                                                                                                                                                            Synthetic
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                                                                                                                                                                                                                                                                                                 AAX77401;
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                                                                                                                                                                                                                                      (hTR). It is used to amplify human staufen (hStau) cDNA synthesised using random hexamer primers and Superscript II reverse transcriptase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Detecting telomerase activity in non-cellular body fluid using a modified
                                                                                                                                                                                    Human staufen polypeptide useful in methods for identifying telomerase
                                                                                                                                                                                                                               The present sequence is a PCR primer specific to human telomerase RNA
                                                                                                                                                                                                                                                                                                 Gaps
       PCR primer; human telomerase RNA; hTR; amplify; human staufen cDNA; hStau; synthesised; random hexamer primer; Superscript II reverse transcriptase; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Telomerase; body fluid; cancer; tumour; screening; TRAP; diagnosis; telomeric repeat amplification protocol; detection; PCR primer; ss.
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                                                                                                                                                                                                                                                                               ch 5.8%; Score 26; DB 1; Length 26; 1 Similarity 100.0%; Pred. No. 59; 26; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                Sequence 26 BP; 7 A; 3 C; 9 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                              Human telomerase RNA template PCR primer F3B.
                                                                                                                                   (UYJO ) UNIV JOHNS HOPKINS SCHOOL MEDICINE
                                                                                                                                                                                                                                                                                                                  145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Highsmith E,
                                                                                                                                                                                                                                                                                                                           CTTCCACCGTTCATTCTAGAGCAAAC 1
                                                                                                                                                                                                              Disclosure; Page 15; 50pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (UYMA-) UNIV MARYLAND BALTIMORE
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                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1999-508655/42.
                                                                                                                                                                    WPI; 1999-620168/53
                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 26; Conserv
                                                                                                                                                    Le S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                 Homo sapiens
                                                                WO9951255-A1
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                                                                                                                                                    Greider CW,
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                                                                                  14-OCT-1999
                                                                                                                                                                                             inhibitors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                             AAZ08703;
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cellular portion of body fluid from a cancer patient using a modified cellular portion of body fluid from a cancer patient using a modified tellular portion of body fluid from a cancer patient using a modified cellular portion of a modified cancer comprises: (a) removing the cellular portion of a body fluid specimen from the patient; (b) preparing a protein extract from the body fluid remainder; (c) assaying the extract for the presence and quantity of telomerase RNA or telomerase activity; and (d) comparing the results with normal levels, to determine the presence of cancer. The methods are therapy effectiveness Unlike prior art telomerase activity assays in cancer patients, the method allows noninvasive sample collection. The methods are also more reliable and less tumour specific than other methods which detect circulating tumour markers. The present sequence represents a human telomerase RNA template PCR primer used in the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Detecting telomerase RNA in urine - useful for diagnosis of bladder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Telomerase; human; diagnosis; bladder cancer; detection; urine;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1; Length 26;
59;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 26 BP; 8 A; 6 C; 7 G; 5 T; 0 U; 0 Other;
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Disclosure; Page 16; 32pp; English
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Best Local Similarity
Matches 26; Conserv
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US09436060A-16.rng.sl

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98WO-US024996.
                                                                              97US-00974180.
pGC6; human; PCR primer; ss.
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                                                                                             GERO-) GERON CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
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                                                                              19-NOV-1997;
                               WO9925878-A2
                                                              19-NOV-1998;
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                                              27-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                         AAX01541;
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                                                                                                                                                                                                                                                                                                                                                                                         RESULT 67
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                                                                                                                                                                                                                                                                                                                                                                  This invention describes a novel method for diagnosing bladder cancer which comprises detecting telomerase RNA in a urine sample. The method of the invention has greater sensitivity and reliability than assays for telomerase activity (cf. WO 975871). This sequence represents a primer used in the method of the invention
                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cellular senescence; modulator; GC6 gene; senescent gene expression;
                                                                                                                                                                                                                                                                                                                           Detecting telomerase RNA in urine - useful for diagnosis of bladder
                                                                                                                                                         Telomerase; human; diagnosis; bladder cancer; detection; urine;
PCR primer; ss.
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100.0%; Pred. No. 59;
iive 0; Mismatches 0; Indels
       DB 1; Length 26; 59;
                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 26 BP; 7 A; 3 C; 9 G; 7 T; 0 U; 0 Other;
        5.8%; Score 26; DB
100.0%; Pred. No. 59;
ive 0; Mismatches
                                                                                                                                               Human telomerase RNA PCR primer TE-hTR3.1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                       79
                                                   1 AACTGAGAAGGGCGTAGGCGCCGTGC 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CTTCCACCGTTCATTCTAGAGCAAAC 1
                                        54 AACTGAGAAGGGCGTAGGCGCCGTGC
                                                                                                                                                                                                                                                                                                                                                     Claim 6; Page 10; 13pp; German.
                                                                                                                                                                                                                                                                              (HOFF ) ROCHE DIAGNOSTICS GMBH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAX77131 standard; DNA; 26 BP
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                                                                                                AAX77402 standard; DNA; 26
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                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                             WPI; 1999-349242/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PCR primer hTR S328
                 Best Local Similarity
                                                                                                                                                                                                 Homo sapiens.
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                         56;
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                                                                                                                                                                                        Synthetic
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                                                                                                                                                                                                                                                                                              Emrich T;
                                                                                                                AAX77402;
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         Query Match
                                                                                                                                                                                                                                                                                                                                        cancer.
                         Matches
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New human GC6 gene, useful for identifying agents for treating diseases and/or conditions associated with cell senescence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TPC2, TPC3; human; telomere length regulation; cancer; pregnancy; fertility; diagnosis; therapy; PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 TTGGGCTCTGTCAGCCGCGGGTCTCT
                                                                                                     Example 5; Page 74; 79pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCR primer for Human TPC3 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP.
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96US-00583808.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAX01541 standard; DNA; 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-APR-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 100.
Matches 26; Conservative
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35

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Gaps

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Indele

100.0%; Prec. ...

ВР

Score 26;

5.8%;

DB 1; Length 26; 59;

G; 7 T; 0 U; 0 Other;

primer;

RT-PCR; PCR 88.

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Detecting telomerase activity in samples, useful for diagnosis of cancer and micrometastasis, comprises treating sample with solid phase, removing solid phase and treating to elute bound telomerase.
                                                                                                                                                                                                                                          Human telomerase RNA reverse transcriptase PCR primer #2.
                                                                                                                                                                                                                                                                  hTR; reverse transcriptase; micrometastasis; diagnosis;
                                                                                145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                        CTTCCACCGTTCATTCTAGAGCAAAC 1
                                                                                                                                                                                                                                                                                                                                                                             01-FEB-2000; 2000WO-IB000100.
                                                                                                                                                                                                                                                                                                                                                                                                  99GB-00002302
           BP; 7 A; 3 C; 9
                                                                                                                                                                   AAA88250 standard; DNA; 26
                                                                                                                                                                                                                  (first entry)
                              Query Match
Best Local Similarity 100.
Matches 26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Larsen F, Skaanseng
                                                                                                                                                                                                                                                                                                                                                                                                                         (LARS/) LARSEN F.
(SKAA/) SKAANSENG M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2000-491281/43
                                                                                                                                                                                                                                                                   telomerase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
les 26; Conserv
                                                                                                                                                                                                                                                                                                                             WO200046601-A1
                                                                                                                                                                                                                                                                                                                                                                                                  32-FEB-1999;
                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                  15-DEC-2000
          Sequence 26
                                                                                                                                                                                                                                                                                                                                                     10-AUG-2000
                                                                                                        56
                                                                                                                                                                                           AAA88250;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                           RESULT 69
                                                                                                                                                         AAA88250,
                                                                                                                                                                                          × 8
                                                                                                      용
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                                                                                                                            This sequence represents a PCR primer for DNA encoding the human TPC3 protein, which is contained within the recombinant mammalian host cell of the invention. The invention provides methods and reagents for regulating telomerae and reagents for regulating telomerae activity in mammalian cells as well as for detecting, diagnosing, and treating related diseases and conditions such as cancer, pregnancy, or fertility in humans and other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This sequence represents a PCR primer for DNA encoding the human TPC3 protein, which is contained within the recombinant mammalian host cell of the invention. The invention provides methods and reagents for regulating telomeraes and reagents for regulating telomeraes activity in mammalian cells as well as for detecting, diagnosing, and treating related diseases and conditions such as cancer, pregnancy, or fertility in humans and other
                                                                                                                                                                                                                                                                              Gaps
                                                                     DNA encoding proteins TPC2 and TPC3 - useful for regulating telomere length or modulating telomerase activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA encoding proteins TPC2 and TPC3 - useful for regulating telomere length or modulating telomerase activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   regulation; cancer; pregnancy; primer; ss.
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                                                                                                                                                                                                                                                    5.8%; Score 26; DB 1; Length 26;
100.0%; Pred. No. 59;
1ve 0; Mismatches 0; Indels
                                                                                                                                                                                                                             Seguence 26 BP; 8 A; 6 C; 7 G; 5 T; 0 U; 0 Other;
                       <u>ن</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Feng J;
                      Feng
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Villeponteau B,
                      Villeponteau B,
                                                                                                                                                                                                                                                                                                      TCTAACCCTAACTGAGAAGGGCGTAG 70
                                                                                                                                                                                                                                                                                                                          TCTAACCCTAACTGAGAGGGCGTAG 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TPC2; TPC3; human; telomere length
fertility; diagnosis; therapy; PCR
                                                                                                                                                                                                                                                      5.5.,
100.0%; Pro
                                                                                                        Example, Col 38; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example; Col 38; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR primer for Human TPC3 gene
                                                                                                                                                                                                                                                                                                                                                                                        BP
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ID AAX01542 standard; DNA; 26
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                                                                                                                                                                                                                                                                              Conservative
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                       Andrews WH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1999-152104/13.
                                              WPI; 1999-152104/13
(GERO-) GERON CORP.
                                                                                                                                                                                                                                                                  Local Similarity
nes 26; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TPC3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-SEP-1996;
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05-JAN-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                      Adams RR,
                                                                                                                                                                                                                                                                                                      45
                                                                                                                                                                                                                                                                                                                                                                                                               AAX01542;
                                                                                                                                                                                                                                                      Query Match
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Matches
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                                                                                                                        activity in a sample. The method comprises treating the sample with a sample of bind telomerase, separating the solid phase from the sample to form a test sample which may be treated to elute bound telomerase and assaying the sample for telomerase activity. Also described are: (1) a kit (II) for detecting telomerase activity, comprising a solid phase and one or more components for assaying telomerase activity, and (2) a component (III) of an assay system for telomerase activity, comprising a solid phase for binding telomerase on which is present a substrate for telomerase elongation. (I) is useful for cancer diagnosis or prognosis and detection of micrometastasis as detection of telomerase activity is indicative of sancer or micrometastasis. The solid phase used in (I) is useful for separating telomerase from a sample and therefore for detecting telomerase from a sample and therefore for detecting and may
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCR primer for
                                                                                       present invention describes a method (I) for detecting telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gapa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     telomerase activity. (II) is useful for detection of cancer cells an also comprises means for assaying an mRNA diagnostic for cancer. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       numan telomerase RNA, which is used in an example from the present
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       present sequence represents a reverse transcriptase (RT)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 26 BP; 7 A; 3 C; 9 G; 7 T; 0 U; 0 Other;
Example 11; Page 38; 68pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.08;
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The present invention describes a method (I) for detecting telomerase activity in a sample. The method comprises treating the sample with a solid phase to bind telomerase, separating the solid phase from the sample to form a test sample which may be treated to elute bound to sample to form a saying the sample for telomerase activity. Also described are: (1) a kit (II) for detecting telomerase activity, in the components for assaying telomerase activity, components for assaying comprising a solid phase and one or more components for assaying telomerase activity, component (III) of an assay system for telomerase activity, comprising a solid phase for binding telomerase activity, comprising a solid phase for binding components on which is present a substrate for telomerase elongation. (I) is useful for cancer or micrometastasis. The solid phase used in (I) is useful for cancer or micrometastasis. The solid phase used in (I) is useful for separating telomerase from a sample and therefore for detecting cancer cells and also comprises means for assaying an mRNA diagnostic for cancer. The present sequence represents a reverse transcriptase (RT) FCR primer for intermity telomerase RNA, which is used in an example from the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Detecting telomerase activity in samples, useful for diagnosis of cancer and micrometastasis, comprises treating sample with solid phase, removing solid phase and treating to elute bound telomerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; telomerase; hTR; reverse transcriptase; RT-PCR; PCR primer; detection; cancer; micrometastasis; diagnosis; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human telomerase RNA reverse transcriptase PCR primer #1.
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CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                        CTTCCACCGTTCATTCTAGAGCAAAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 11; Page 38; 68pp; English.
                                                                                                                                                                                                                                                                        ВР
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                                                                                                                                                                                                                                                                        AAA88249 standard; DNA; 26
                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (LARS/) LARSEN F.
(SKAA/) SKAANSENG M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-491281/43.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      invention
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                                                                                                                                                                                                                                                                                                                                                    AAA88249;
        145
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                                                                                                                                                                                                    RESULT 70
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The invention relates to detecting human telomerase RNA template (HTR) RNA can a bodily fluid, comprising amplitying RNA extracted from plasma or in a bodily fluid, comprising amplitying RNA extracted from plasma or serum sample, or its corresponding CDNA comprising(1), using primers or probes that target (1) or CDNA and detecting qualitatively or consequence of (1) or cDNA product. The method is useful for detecting (1) in a bodily fluid, which is useful for consequence of (1) or cDNA product. The method is useful for detecting (1) in a bodily fluid, which is useful for consequence of (1) or cDNA product. The method is useful for detecting (1) in a bodily fluid, which is useful for consequence of (1) or cDNA product. The method is useful for detecting of (1) expressing cells or tissue which include a malignancy preferably a cancer of breast, ovarian, head and neck, lung, cervical, colorectal, gastric, liver, pancreatic, bladder, prostate, consequence of cervical intracapithalial neophasia, bronchial dysplasia, atypical carricraphical in repeats of arcombar and an estur carcinoma in situ, colorectal adenoma, atypical endometrial hyperplasia, or Barrett's ossophagus, where the carpina of the breast, ductal carcinoma in situ. The method is a consequence of therapy, which comprises selecting the human for the therapy after the also useful for treating a human with cancer for telomerase-directed therapy, which comprises selecting the human for the therapy after detection of (1), for determining a need for diagnostic test in a human. Consequence of (1), for determining a need for diagnostic test in a human (1) provides a marker which is utilised as a diministered to a human. (1) provides a marker which is utilised as a prognosis in these patients. The method also allows is dentification or analysis, either quantitatively or qualitatively, of (1) in plasma or premalignant or malignant cancer for longered therapy as an early consequence of for further therapy or who has completed therapy or treatment failure.
                                                                                                                                                                                                                            cancer; breast; ovarian; head and neck; lung; cervical; colorectal; gastric; liver; pancreatic; bladder; prostate; brain; kidney; oesophagus; melanoma; sarcoma; premalignancy; carcinoma in-situ; cervical dysplasia; bronchial dysplasia; cervical intraepithelial neoplasia;
                                                                                                                                                                                                        telomerase-associated RNA template; hTR; endometrial; malignancy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Detecting human telomerase RNA template RNA or human telomerase reverse transcriptase protein RNA in bodily fluid, useful as marker for diagnosing, monitoring or treating cancer, carcinoma in situ or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to detecting human telomerase RNA template (hTR)
                                                                                                                                                          Human telomerase-associated RNA template (hTR), PCR primer hTR2.
                                                                                                                                                                                                                                                                                                                                 atypical hyperplasia; colorectal adenoma;
atypical endometrial hyperplasia; tumour; Barrett's oesophagus;
telomerase-directed therapy; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Page 14; 30pp; English.
            ABK48024 standard; DNA; 26 BP.
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                                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ONCO-) ONCOMEDX INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                  18-JUN-2002
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                                                                    ABK48024;
ABK48024/c
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Gaps

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TCTAACCCTAACTGAGAAGGGCGTAG

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Best Local Similarity 100. Matches 26; Conservative

TCTAACCCTAACTGAGAAGGGCGTAG 26

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cancer; breast; ovarian; haad and neck; lung; cervical; colorectal; gastric; liver; pancreatic; bladder; prostate; brain; kidney; oesophagus; melanoma; sarcoma; premalignancy; carcinoma in-situ; cervical dysplasia; bronchial dysplasia; cervical intraepithelial neoplasia; atypical hyperplasia; colorectal adenoma; atypical endometrial hyperplasia; tumour; Barrett's oesophagus;
method allows the development and application of telomerase-specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; telomerase-associated RNA template; hTR; endometrial; malignancy;
                     therapy even when only premalignant tumours, early cancer, or occult cancer or metaetcasis auch as following resection or in minimal residual disease are present. The present sequence represents a PCR primer for human telomerase-associated RNA template (hTR)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  reverse
                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human telomerase-associated RNA template (hTR), PCR primer hTR1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Detecting human telomerase RNA template RNA or human telomerase transcriptase protein RNA in bodily fluid, useful as marker for diagnosing, monitoring or treating cancer, carcinoma in situ or
                                                                                                                                                                                                        ..
                                                                                                                                                           5.8%; Score 26; DB 1; Length 26; 100.0%; Pred. No. 59;
                                                                                                                                                                                                        0; Indels
                                                                                                                       Sequence 26 BP; 7 A; 3 C; 9 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      telomerase-directed therapy; primer; ss.
                                                                                                                                                                                                                                            145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                    26 CTTCCACCGTTCATTCTAGAGCAAAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Page 14; 30pp; English.
                                                                                                                                                                                                                                                                                                                                                                                           ABK48023 standard; DNA; 26 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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Best Local Similarity 100.0
Matches 26; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                    ABK48023;
                                                                                                                                                                                                                                                                                                                                                     RESULT 72
                                                                                                                                                                                                                                                                                                                                                                        ABK48023
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the atypical endometrial hyperplasia, or Barrett's oesophagus, where the human is at risk for developing a malignancy or premalignancy or is known to have malignancy, premalignancy or carcinoma in situ. The method is also useful for treating a human with cancer for telomerase-directed therapy, which comprises selecting the human for the therapy after ceferction of [1], for determining a need for dagnostic test in a human with malignancy or premalignancy amonitoring a therapy of the manignancy or premalignancy and for monitoring a therapy of with malignancy or premalignancy and for monitoring a therapy of computer therapeutic effect has been achieved, or whether additional or more advanced therapy is required, and to assess prognosis in these patients. The method also allows identification or whether additional or more advanced therapy is required, and to assess prognosis, either quantitatively or qualitatively, of [1] in plasma or canalysis, either quantitatively or qualitatively, of [1] in plasma or canalysis, either quantitatively or qualitatively, of [1] in plasma or canalysis, either quantitatively or qualitatively, of [1] in plasma or canalysis, either quantitatively or qualitatively, of [1] in plasma or canalysis, either quantitatively or premalignant lesions, and thus allow stratification of such core their risk of residual cancer following surgery, and their consect for further therapy or who has completed therapy as an early or hear who has completed therapy as an early or them man and the planes of the planes of development and application of telomerase specific themselved to the planes of the planes of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                           therapy even when only premalignant tumours, early cancer, or occult cancer or metastasis such as following resection or in minimal residual disease are present. The present sequence represents a PCR primer for human telomerase-associated RNA template (hTR)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Peptide products of the human TPC2 and TPC3 gene are involved in regulation of telomere length and activity are useful to diagnose and treat telomere length and activity-related diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to methods and reagents for regulating telomere length and for modulating telomerase activity in mammalian cells. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TR; telomerase activity-related disease; therapy; fertility; RT-PCR primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human telomerase (hTR) cDNA amplifying R3c downstream RT-PCR primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1; Length 26; 59;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 26 BP; 8 A; 6 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Adams RR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Pred. No. 59; cive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5.8%; Score 26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TCTAACCCTAACTGAGAAGGGCGTAG 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              rcraacccraacrcagaagggggrag 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Andrews WH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example, Col 38; 60pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       95US-0003492P.
96US-00583808.
96US-00710249.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-MAR-2002 (first entry)
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Best Local Similarity 100.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Villeponteau B, Feng J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-033174/04.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; telomerase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cancer; pregnancy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GERO-) GERON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-SEP-1995;
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invention also relates to purified, synthetic or recombinant peptides such as TPC2 or TPC3 used for detecting regulators of telomere length and telomerase activity in mammalian cells and for a variety of related diagnostic and therapeutic purposes. The method is useful for screening, diagnosing, monitoring and treating diseases and other conditions such as ancer, pregnancy, fertility, telomere length and telomerase-activity. The present sequence is a reverse transcription (RT) PCR primer used for amplifying human telomerase (hTR) cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to methods and reagents for regulating telomere length and for modulating telomerase activity in mammalian cells. The invention also relates to purified, synthetic or recombinant peptides such as TPC2 or TPC3 used for detecting regulators of telomera length and telomerase activity in mammalian cells and for a variety of related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Peptide products of the human TPC2 and TPC3 gene are involved in regulation of telomere length and activity are useful to diagnose and treat telomere length and activity-related diseases.
                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; telomerase; TR; telomerase activity-related disease; therapy; cancer; pregnancy; fertility; RT-PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human telomerase (hTR) cDNA amplifying F3b upstream RT-PCR primer.
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                                                                                                                                                                                                   Score 26; DB 1; Length 26;
Pred. No. 59;
                                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                   Sequence 26 BP; 7 A; 3 C; 9 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 26 BP; 8 A; 6 C; 7 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                              100.0%; Pred. ...
                                                                                                                                                                                                                                                                             145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                   CTTCCACCGTTCATTCTAGAGCAAAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Andrews WH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example; Col 38; 60pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           95US-0003492P.
96US-00583808.
96US-00710249.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                          26; Conservative
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                                                                                                                                                                                                                         Best Local Similarity
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05-JAN-1996;
13-SEP-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
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                                                                                                                                                                                                                                                                                                              56
                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                          Matches
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Accurate, reliable diagnosis and/or prognosis of cancer, e.g. breast cancer, by analyzing the RNA components of telomerase in plasma or serum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      monitoring the evolution of cancers. The method comprises analysing enzyme telomerase RNA in blood plasma or serum. The method is typically used for diagnosing breast, ovarian, stomach or colon cancer and/or monitoring the evolution of the cancers after treatment by chemotherapy or operations. The present sequence is a PCR primer for human telomerape RNA (HTR), which was used in the example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; ss; PCR; telomerase; cancer; pregnancy; fertility; neoplasm; hTR; RT-PCR; reverse transcriptase PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                           telomerase RNA; PCR primer; cancer; breast; ovarian; stomach; hTR; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to a method for diagnosing and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
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59;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 26 BP; 3 A; 6 C; 11 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human telomerase gene, hTR, RT-PCR primer #2.
                                                                                                                                                                       Human telomerase RNA, hTR, sense PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Prec. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 26;
Pred. No.
70
                       1 rcraacccraacrgagaagggcgrag 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 60 GAAGGGCGTAGGCGCCGTGCTTTTGC 85
45 TCTAACCCTAACTGAGAAGGGCGTAG
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                                                                                         BP
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                                                                                                                                                                                                                                                                                                                     26-MAY-2000; 2000EP-00111370
                                                                                                                                                                                                                                                                                                                                              26-MAY-2000; 2000EP-00111370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example; Col 3; 6pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                           Stroun M, Anker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADG82593/c
ID ADG82593 standard; DNA; 26
                                                                                         ABA95496 standard; DNA; 26
                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                      (CHEN/) CHEN X Q. (STRO/) STROUN M.
                                                                                                                                                                                                                                                                                                                                                                                                 (ANKE/) ANKER P.
                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                  EP1158055-A1.
                                                                                                                                            12-MAR-2002
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                                                                                                                   ABA95496;
                                                                                                                                                                                                                                                                                                                                                                                                                           Chen XQ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADG82593;
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                                                                                                                                                                                               Human;
                                                                                                                                                                                                                colon;
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                                                                             ABA95496
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Gaps

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5.8%; Score 26; DB 1; Length 26; 100.0%; Pred. No. 59; iive 0; Mismatches 0; Indels

Conservative

26;

Matches

Query Match Best Local Similarity

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The invention relates to a monoclonal or isolated polyclonal antibody specifically binds human TPC2 or TPC3 (not defined, appearing as ADG82569 and ADG82571). Also included are an antibody obtained by collecting attribution a subject immunised with a peptide comprising at least 10 cantiguous amino acids of TPC-2 pr TPC-3, a host cell secreting the cartibody, a composition for obtaining a TPC2-specific antibody comprising at least 10 contiguous amino acids ADG82569, a composition for obtaining at least 10 contiguous amino acids ADG82569, a composition for obtaining a TPC3-specific antibody comprising at least 10 contiguous amino acids of ADG82571, an isolated, recombinant or synthetic nucleic acid encoding a peptide immunogenic for TPC2-specific antibody or TPC3-antibody and a host cell containing the nucleic acid. The antibody is useful for determining a condition in a subject associated with a high level of TPC2 or TPC3, and is useful for screening, diagnosing and monitoring conditions auch as secilated with inappropriate expression of TPC2 or TPC3 such as neoplasia. The present sequence is a reverse transcriptase (RT)-PCR primer used to analyse the expression profile of the human telomerase gene, which is co-expressed with TPC-2 and TPC-3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel monoclonal or isolated polyclonal antibody that specifically binds human TPC2 or TPC3. useful for treating neoplasia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; gene therapy; thymidine kinase gene; anticancer therapy; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human telomerase RNA gene (hTR) specific primer TRC3R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 26 BP; 8 A; 6 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                  Andrews WH, Adams RR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5.8%; Score 26; DB
100.0%; Pred. No. 59;
:ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example D; SEQ ID NO 25; 62pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           45 TCTAACCCTAACTGAGAGGGCGTAG 70
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96US-00583808.
96US-00710249.
98US-00220157.
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                                                                                         29-JUN-2001; 2001US-00895606
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                                                                                                                                                                                                                                                                                                                                                                  Feng J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 100.
Matches 26; Conservative
                                                                                                                                                                                                                                                 VILLEPONTEAU B.
                                                                                                                                                                                                                                                                        FENG J.
ANDREWS W H.
                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-051519/05.
                                                                                                                                                                                                                                                                                                                     (ADAM/) ADAMS R R
                                                                                                                                                                                                                                                                                                                                                               Villeponteau B,
US2003207404-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene therapy;
                                                                                                                                09-SEP-1995;
05-JAN-1996;
13-SEP-1996;
                                                                                                                                                                                                       23-DEC-1998;
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                                              06-NOV-2003
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                                                                                                                                                                                                                                                 (VILL/)
                                                                                                                                                                                                                                                                        (FENG/)
                                                                                                                                                                                                                                                                                                  (ANDR/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a monoclonal or isolated polyclonal antibody specifically binds human TPC2 or TPC3 (not defined, appearing as ADG8256) and ADG82571). Also included are an antibody obtained by collecting a natiserum from a subject immunised with a peptide comprising at least 10 contiguous amino acids of TPC-2 pr TPC-3, a host cell secreting the ar least 10 contiguous amino acids of TPC-2 pr TPC-3, a host cell secreting the a least 10 contiguous amino acids of ADG82569, a composition for obtaining a TPC3-specific antibody comprising a TPC3-specific antibody comprising a TPC3-specific antibody contiguous amino acids of ADG82571, an isolated, recombinant or synthetic nucleic acid encoding a peptide immunogenic for TPC2 sepecific antibody or TPC3-antibody and a host cell containing the nucleic acid. The antibody is useful for ceremining a condition in a subject associated with a high level of TPC2 or TPC3, and is useful for screening, diagnosing and monitoring conditions such as cancer, pregnancy or fertility. The antibody is also useful for treating conditions associated with inappropriate expression of TPC2 or TPC3 such as neoplasia. The present sequence is a reverse transcriptes (RT)-PCR primer used to analyse the expression profile of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; 88; PCR; telomerase; cancer; pregnancy; fertility; neoplasm; hTR; RT-PCR; reverse transcriptase PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel monoclonal or isolated polyclonal antibody that specifically binds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               human telomerase gene, which is co-expressed with TPC-2 and TPC-3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5.8%; Score 26; DB 1; Length 26; 100.0%; Pred. No. 59;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 human TPC2 or TPC3. useful for treating neoplasia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 26 BP; 7 A; 3 C; 9 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Andrews WH, Adams RR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Preq. ....
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human telomerase gene, hTR, RT-PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example D; SEQ ID NO 26; 62pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         95US-0003492P.
96US-00583808.
96US-00710249.
98US-00220157.
                                                                                                                                                          2001US-00895606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADG82592 standard; DNA; 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                 Villeponteau B, Feng J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                     VILLEPONTEAU
                                                                                                                                                                                                                                                                                                                                                               ANDREWS W H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2004-051519/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                       ADAMS R R.
                                                                 US2003207404-A1.
                       Homo sapiens.
                                                                                                                                                        29-JUN-2001;
                                                                                                                                                                                                                                                                                                                                          FENG
                                                                                                                                                                                                                          05-JAN-1996;
13-SEP-1996;
                                                                                                                                                                                                                                                                        23-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11-MAR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26;
                                                                                                              06-NOV-2003
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                                                                                                                                                                                                                                                                                                                (VILL/) (FENG/) (ANDR/) (ADAM/) (ADAM/)
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Gaps

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DB 1; Length 26; 59; 0; Indels

Matches

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ADG82592

Corey DR;

Norton JC,

Shay JW,

Piatyszek MA,

96US-00630019.

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(TEXA ) UNIV TEXAS SYSTEM.
09-APR-1996;
                                                  Wright WE,
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                                                                                                                                                                                                                                          RNA (TR) component genes. The TR gene promoter can be linked to a heterologous gene, especially a gene encoding a cytotoxin, for therapy of cancer, especially neoplasias. The telomerase is necessary for the cancer, especially neoplasias. The telomerase is necessary for the curestricted proliferative capacity of many human cancers. Mutation or upregulation of the telomerase repression pathway may cause reactivation or upregulation of telomerase expression in cancer. Substances, clamping the methods, can be used to block transcription from the TR gene promoter through interaction of the 5' regulatory sequences. These cubstances, e.g. antitaense oligonuclectides, transcription factors, peptide nucleic acids and factors that disrupt signal transduction, are useful for cancer therapy. In particular, gene therapy vectors cancer therapy. In particular, gene therapy vectors cancer therapy. In particular, gene therapy vectors can be used to convert a producy, e.g. gancyclovir, so that neoplasia can be controlled or treated. Direct down-regulation of telomerase RNA gene through manipulation of transcription factors may be effective anticancer therapy and the cloning of the hTR gene promoter activity. Sequences AAZ07623-80 represents PCR primers for amplifying
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                                                                                                                                                                                                                                  The invention relates to promoter regions from mouse and human telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Peptide nucleic acid; PNA; telomerase; ribonucleoprotein enzyme; cancer; inhibitor; neoplasia; neurodegenerative disease; aging; hyperplasia; AIDS; HIV; fungal infection; forensic identification; detect; tumour; paternity testing; ss.
                                                                                                                                                                    Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 25 BP; 6 A; 7 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5.5%; Scor.
100.0%; Pred. No. vc.
'-^ 0; Mismatches
                                                                                         (CANC-) CANCER RES CAMPAIGN TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              human TR gene (hTR) promoter sequence
                                                                                                                                                                                                        Disclosure; Fig 6; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Telomerase RNA target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAA37582 standard; RNA; 25 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   97US-00838545.
                                                                98GB-00001902.
                                      99WO-GB000308
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 100.0
                                                                                                                                          WPI; 1999-479183/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-APR-1997;
                                      29-JAN-1999;
                                                                29-JAN-1998;
                                                                                                                                                                                gene therapy
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              05-AUG-1999.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA37582;
                                                                                                                 Keith WN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 79
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The present sequence represents a human telomerase RNA sequence which is the target of peptide nucleic acid molecules (PNA). The PNA molecules the target of peptide nucleic acid molecules (PNA). The PNA molecules contained within the RNA component of mammalian telomerase, and inhibit telomerase activity. Telomerase is a ribonucleoprotein enzyme that telomerase activity. Telomerase is a ribonucleoprotein enzyme that telomerase activity. Telomerase is a ribonucleoprotein enzyme that contained within the RNA component of the enzyme. The nucleoride sequence contained within the RNA component of the enzyme. The nucleoride sequence contained within the RNA component of the enzyme. The nucleoride sequence contained within the RNA component of the PNA bases which include the sequence GTTAGG. The uncharged nature of the PNA bases the melting temperature of associating strands, increases the rate of association by the targeted nucleic acids, and afformed grader resistance of degradation by proceases or nucleases. The terapeutic PNAs may be used for treating disease conditions such as cancers, neoplasia, hyperplasia, therefore metabolism or telomerase diseases characterized by abnormal telomera metabolism or telomerase characterized by abnormal telomera metabolism or telomerase deformedeformel and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization probes to detect or quantitate polymucleotides having a human telomerase RNA (hTR) sequence. PNA probes are also used for forensic identification of individuals, e.g. paternity testing, based on hTR gene restriction of individuals, e.g. paternity testing, based on hTR gene restriction almost and elect the RNA component of a mammalian telomerase and as inhibitors of telomerase activity. The method of the present invention can be used to detect an immortal or meoplastic cell or tumour issue or cancer of any origin, provided the meaning activity and its RNA component cells any detect an immortal or meoplastic cell expresses telome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ò
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                                                                     New peptide nucleic acid (PNA) compounds that inhibit telomerase activity in mammalian cells is useful as probes to detect the RNA component of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     41 TTTGTCTAACCCTAACTGAGAAGGG 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human telomerase RNA (hTR) strand #1.
                                                                                                                                                                                                                  Example 1; Col 27-28; 45pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAS15453 standard; RNA; 25 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14-FEB-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity 72.0
les 18; Conservative
                                                                                                                                                 mammalian telomerase.
WPI; 2000-292432/25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAS15453;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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Matches
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05-JUN-2000; 2000WO-US015544.
                                                                                 (AUJL/)
(WIEN/)
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                                                                                                                                              δu JL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAV41169,
  à
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to peptide nucleic acids (PNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in mammalian cells by hybridising to the RNA component of mammalian telomerase. The PNAs are useful as probes to detect the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polymucleotide having the human telomerase RNA component (hTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the HTR gene RRLP pattern. The PNA can be further used for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmaceuticals (such as antineoplasis, human immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomerase detabolism or telomerase activity. The present sequence representing hTR RNA strand #1 is used to the the inhibition of telomerase activity by the PNAs of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                               New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases.
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                                                                                                                                                                                                                                                                         Norton JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1; Length 25;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 25 BP; 7 A; 5 C; 6 G; 0 T; 7 U; 0 Other;
                                                                                                                                                                                                                                                                         Corey DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pred. No. 68;
7; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 25;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   65
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                                                                                                                                                                                                                                                                         Piatyszek MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 41 TTTGTCTAACCCTAACTGAGAAGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Col 29; 46pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human telomerase PCR primer #1
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                                                                                                                          99US-00349532.
                                                                                                                                                                96US-00630019.
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                                                                                                                                                                                                                             (TEXA ) UNIV TEXAS SYSTEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18; Conservative
                                                                                                                                                                                                                                                                       Shay JW, Wright WE,
                                                                                                                                                                                                                                                                                                                 WPI; 2001-638024/73
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  Homo sapiens
                                         US6294650-B1
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                                                                                                                        08-JUL-1999;
                                                                                                                                                                09-APR-1996;
                                                                                                                                                                                        09-APR-1997;
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                                                                               25-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              invention
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Best Local S
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80 X C C C C C C C C C C C C C X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B X B
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                                                                                                                                                                                                                                                                                                                                                                                            The present invention provides a method for inhibiting or reducing the growth of a cell which involves administering to the cell a telomere damage inducing agent and a telomerase inhibitory agent. This can be used in the treatment of aberrant cell growth, including cancers
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                                                                                                                                                                                                                                  Inhibiting or reducing growth of cell for treating cancer, comprising administering telomere damage-inducing agent and telomerase inhibitory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RNA component; human telomerase; antisense oligonucleotide; infection; neuroblastoma; bladder cancer; colon cancer; prostate cancer; cancer; contraception; sterilisation; immunosuppression; therapeutic; hTR; immune system down-regulation; anti-inflammatory therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RNA component of human telomerase (hTR) antisense oligo 14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1; Length 25;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Seguence 25 BP; 4 A; 5 C; 4 G; 12 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5.5%; Score 25; 00.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              161 TAGAGCAAACAAAAATGTCAGCTG 185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TAGAGCAAACAAAAATGTCAGCTG 1
                                                                                                                                                                                                                                                                                                                                             Sxample 7; Page 62; 97pp; English.
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99US-0137549P
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96US-00770565
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity 100.
Matches 25; Conservative
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                                                                                                                                                                                   WPI; 2001-071022/08
                                                 AU J L.
WIENTJES G.
                                                                                                                              Wientjes G;
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                                                                                                                                                                                                                                                                                            agent to the cell.
04-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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antisense oligonucleotides to the RNA component of human telomerase (hTR). These antisense oligonucleotides specifically hybridise to a nucleotide component of the hTR, but that does not hybridise to a sequence within the remplate region of hTR. These oligonucleotides may specifically be used for detection of an RNA component of human telomerase in a sample. This is useful for diagnosing cancer (especially neuroblastoma, bladder, colon and prostate cancer), and providing prognosis for a cancer patient. The inhibitory coligonucleotides can inhibit the telomerase activity level in a cell by interfering with transcription of the RNA component, decreasing the half-coller of the telomerase RNA component transcript, inhibiting assembly of the RNA component transcript, inhibiting the coller of the RNA component into the telomerase Answer of inhibiting the coller in vivo. They can be used in therapeutics for treating or cells in vivo. They can be used in therapeutics for treating or preventing cancer, for contraception or sterilisation, for immunosuppression, and for selectively down-regulating specific branches confine immune system, e.g. a specific subset of rcells, in anti-cinflammatory therapies or for treating infections by, e.g. yeast,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This is an oligonucleotide used in the measurement of Tm values and their complimentary peptide nucleic acids (PNAs), (e.g. AAT89228-T89233, AAT89243-T89244). PNAs hybridise specifically to an RNA component of mammalian telomerase, and include the sequence GGG for specific hybridisation to the template region of this component. PNAs can be used as probes to detect the RNA component of mammalian telomerase and as
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Peptide nucleic acid, PNA; cancer, telomerase, probe, hybridisation, inhibitor; human telomerase RNA; hTR; PCR; oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New peptide nucleic acids hybridising to mammalian telomerase RNA - to inhibit telomerase, for treating tumours and other proliferative diseases, also for diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA oligonucleotide 1, used in the measurement of Tm values.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1; Length 30;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 30 BP; 4 A; 10 C; 8 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     5.4%; Score 24.4; D
96.2%; Pred. No. 97;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CCTTTCAGGCCGCAGGAAGAGGAACG 380
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 2; Page 49; 76pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 97WO-US005931.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1997-512647/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-APR-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        telomerase. Detecting telomerase allows early diagnosis of tumour or infection, before clinical signs manifest. Telomerase inhibitors directed against e.g. Trypanosoma should cause fewer side effects than drugs currently used to treat such infections. The DNA encodes those parts of MTR RNA essential for activity but are significantly shorter than the
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cancer; eukaryotic parasite; hTR; vertebrate telomerase; yeast; protozoa;
inhibitors of telomerase activity, especially in the treatment of cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA encoding essential RNA components of human telomerase - also truncated or recombinant telomerase, useful for diagnosis and treatment of cancer and infection by eukaryotic parasites.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  or to prevent immortalisation of cells in tumours or parasites. The DNA is also used to produce recombinant telomerase, which can then be used conventionally to raise antibodies for diagnostic detection of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                           DB 1; Length 24; 78;
                                                                                                                    0; Indels
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                                     Sequence 24 BP; 7 A; 5 C; 5 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 tumour; antibody; polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                      Human telomerase template PCR primer TRC31.
                                                                                                  Local Similarity 100.0%; Pred. No. 78; es 24; Conservative 0; Mismatches
                                                                               Score 24;
                                                                                                                                                            41 TITGICIAACCCIAACTGAGAAGG 64
                                                                                                                                                                                               1 Trrgrcraacccraacrgagagg 24
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                                                                                                                                                                                                                                                                                                      BP.
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                                                                                 5.3%;
                                                                                                                                                                                                                                                                                                 AAT58807 standard; DNA; 24
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                                                                                                                                                                                                                                                                                                                                             AAT58807;
                                                                                 Query Match
                                                                                                      Best Loca
Matches
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                                                                                                                                                                                                                                                            RESULT 84
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primers AAV68464-65 are used for the RT-PCR amplification of the RNA component of human telomerase (hTR). The invention relates to an activator-antisense complex (A1) that comprises: (a) an antisense oligonucleotide, complementary to a 12-25 nucleotide portion of hTR, with a hydroxyl moiety at the first end; and (b) a linker attached to the first. The activator-antisense complex may be used for inhibiting the growth of a celomerase-expressing malignant cell or tumour. The complex is used to specifically y cleave the ribonucleotide portion of a telomerase enzyme. The complex inhibite growth of telomerase expressing malignant glioma, breast tumour, renal cell cancer. The complex inhibite growth of telomerase expressing malignant glioma, breast tumour, renal cancer.

The complex inhibite growth of telomerase expressing malignant cells from brain tumour malignant glioma, breast tumour, renal call cancer.

C disorders, may be treated including various acute and chronic leukemias, polychemia vera, Hodgkin's and non-Hodgkin's lymphomas, multiple mylenen, Maldenstrom's macroglobulinemia, heavy chain disease, and solid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cleave; brain; tumour malignant glioma; breast tumour; renal cell câncer; melanoma; prostete cancer; leukemia; polytchemia vera; myeloma; sarcoma; Hodgkin's lymphoma; Waldenstrom's macroglobulinemia; heavy chain disease; carcinoma; chemotherapeutic; antisense; RT-PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; telomerase; hTR; activator-antisense complex; malignant; enzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New RNase L activator-telomerase antisense complex - useful to inhibit telomerase activity in telomerase-expressing malignancies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tumours, including numerous sarcomas and carcinomas. The complex is preferably administered in combination with a chemotherapeutic agent, particularly either cisplatin, doxorubicin, mitomycin, daunorubicin, bleomycin, actinomycin D, or neocarzinostatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Torrence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human telomerase RNA (hTR) amplifying RT-PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 24 BP; 3 A; 6 C; 10 G; 5 T; 0 U; 0 Other;
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CTCTGTCAGCCGCGGTCTCTCGG 334
                                                                         24 crcrercaeceeceeercrees 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example; Page 41; 81pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLINIC FOUND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (USSH ) US NAT INST OF HEALTH
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                                                                                                                                                                                                                                                                                       AAV68465 standard; DNA; 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Silverman RH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ношо варіепв
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                                                                                                                                                                                                                                                                                                                                                                         AAV68465;
311
                                                                                                                                                                                                          RESULT 85
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primers AAV68464-65 are used for the RT-PCR amplification of the RNA component of human telomerase (hTR). The invention relates to an activator-antisense complex that comprises: (a) an antisense of longular complementary to a 12-25 nucleotide portion of hTR, with a hydroxyl moiety at the first end; and (b) a linker attached to the first end; and (c) an activator of RNase L attached to the linker. The cutvator-antisense complex may be used for inhibiting the growth of a telomerase-expressing malignant cell or tumour. The complex is used to specifically cleave the ribonucleotide portion of a telomerase enzyme. The complex inhibits growth of telomerase expressing malignant cells from rain tumour malignant glioma, breast tumour, renal cell cancer. Menson therefore and chronic leukemias, collected and prostate cancer. Many other malignancies and related disorders, may be treated including various acute and chronic leukemias, polycthemia vera, Hodgkin's and non-Hodgkin's lay lymphomas, multiple myeloma, waldenstrom's macroglobulinemia, heavy chain disease, and solid tumours, including numerous sarcomas and carcinomas. The complex is
                                                                                                                                                                                                                                                             Human; telomerase; hTR; activator-antisense complex; malignant; enzyme; cleave; brain; tumour malignant glioma; breast tumour; renal cell cancer; melanoma; prostate cancer; leukemia; polycthemia vera; myeloma; sarcoma; Hodgkin's lymphoma; Waldenstrom's macroglobulinemia; heavy chain disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New RNase L activator-telomerase antisense complex - useful to inhibit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      particularly either cisplatin, doxorubicin, mitomycin, daunorubicin,
bleomycin, actinomycin D, or neocarzinostatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     preferably administered in combination with a chemotherapeutic
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                                                                                                                                                                                                                                                                                                                                           carcinoma; chemotherapeutic; antisense; RT-PCR; primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             telomerase activity in telomerase-expressing malignancies.
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                                                                                                                                                                                                                            Human telomerase RNA (hTR) amplifying RT-PCR primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24
CGTGCACCCAGACTCGGCTCACA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  41 TTTGTCTAACCCTAACTGAGAAGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (CLEV-) CLEVELAND CLINIC FOUND. (USSH ) US NAT INST OF HEALTH.
                                                                                                                 ВР
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98US-00018125.
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                                                                                                                 AAV68464 standard; DNA; 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1998-609972/51.
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                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
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03-FEB-1998;
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                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                    AAV68464;
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24; Conservative

Similarity

Query Match Best Local 8

Matches

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Gaps ö

0; Indels

Score 24; DB 1; Length 24; Pred. No. 78;

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Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; gene therapy; thymidine kinase gene; anticancer therapy; human;

PCR primer; ss.

Homo sapiens

Synthetic.

W09938964-A2 05-AUG-1999

Human telomerase RNA gene (hTR) specific primer hTR5.

22-OCT-1999 (first entry)

AAZ07263;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention trained to promoter regions trong and inheritor trained to a cancer, especially a gene promoter can be linked to a cancer, especially neoplasias. The telemerase is necessary for the cancer, especially neoplasias. The telemerase is necessary for the unrestricted proliferative capacity of many human cancers. Mutation or dysregulation of the telemerase repression pathway may cause reactivation or dysregulation of the telemerase expression in cancer. Substances, identified in the methods, can be used to block transcription from the TR cance promoter through interaction of the 5' regulatory sequences. These substances, e.g. antisense oligomucleotides, transcription factors, captule nucleic acids and factors that disrupt signal transduction, are useful for cancer therapy. In particular, gene therapy vectors (especially pGT62-codAupp) comprising the promoter and a viral thymidine cuseful for cancer therapy in particular, gene therapy vectors can be used to convert a prodrug, e.g. gancyclovir, so that neoplasia can be controlled or treated. Direct down-regulation of telemerase RNA gene through manipulation of transcription factors may be effective anticancer therapy and the cloning of the HTM gene promoter allows the analysis of therapeutic molecules which modulate hTM promoter activity. Sequences AAZ07623-80 represents PCR primers for amplifying thuman TR gene (hTR) promoter sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                                                                                                                                                                Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; gene therapy; thymidine kinase gene; anticancer therapy; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5.3%; Score 24; DB 1; Length 24;
100.0%; Pred. No. 78;
iive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                  Human telomerase RNA gene (hTR) specific primer hTR5.
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                                                                                                        AAZ07273 standard; DNA; 24 BP.
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Best Local Similarity 100.
Matches 24; Conservative
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                                                                                                                                                                                                                                                                                                                                                                             PCR primer; 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
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                                                                                       AAZ07273/c
                                                            RESULT 87
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Mouse and human telomerase RNA gene promoters, useful for tumor specific

WPI; 1999-479183/40.

Keith WN;

Disclosure; Fig 6; 109pp; English

(CANC-) CANCER RES CAMPAIGN TECHNOLOGY.

99WO-GB000308 98GB-00001902

29-JAN-1999; 29-JAN-1998;

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RIA (TR) component genee. The TR gene promoter can be linked to a cancer, especially a gene encoding a cytotoxin, for therapy of cancer, especially neoplasias. The telomerase is necessary for the unrestricted proliferative capacity of many human cancers. Mutation or dysregulation of the telomerase repression pathway may cause reactivation or upregulation of telomerase expression in cancer. Substances, identified in the methods, can be used to block transcription from the TR gene promoter through interaction of the 5' regulatory sequences. These substances, e.g. antisense oligonucleotides, transcription factors, peptide nucleic acids and factors that disrupt signal transduction, are useful for cancer therapy. In particular, gene therapy vectors in particular, gene therapy vectors tinase gene can be used to convert a product and a viral thymidine compasia can be controlled or treated. Direct down-regulation of telomerase RNA gene through manipulation of transcription factors may be recommended.
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invention relates to promoter regions from mouse and human telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          effective anticancer therapy and the cloning of the hTR gene promoter allows the analysis of therapeutic molecules which modulate hTR promoter activity. Sequences AAZ07623-80 represents PCR primers for amplifying human TR gene (HTR) promoter sequence
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Best Local Similarity
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Gaps

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46 CTAACCCTAACTGAGAAGGGCGTA 69 CTAACCCTAACTGAGAAGGGCGTA 1

24

ð g AAZ07263/c ID AAZ07263 standard; DNA; 24 BP.

RESULT 88

88

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paternity testing;
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                                                                                                                                                                                                                                                                                                                                                            The invention relates to promoter regions from mouse and human telomerase RNA (TR) component genes. The TR gene promoter can be linked to a neterologous gene, sepecially a gene encoding a cytocain, for therapy of cancer, especially neoplasias. The telomerase is necessary for the unrestricted proliferative capacity of many human cancers. Mutation or corresponding to telomerase repression pathway may cause reactivation of telomerase expression pathway may cause reactivation or dentified in the methods, can be used to block transcription from the TR gene promoter through interaction of the 5' regulatory sequences. These substances, e.g. antisense oligonucleotides, transcription factors, peptide nucleic acids and factors that disrupt signal transduction, are useful for cancer therapy. In particular, gene therapy vectors (sepecially pGTG2-codAupp) comprising the promoter and a viral thymidine kinase gene can be used to convert a producer and a viral thymidine coplasia can be controlled or treated. Direct down-regulation of telomerase RNA gene through manipulation of transcription factors may be telective anticancer therapy and the cloning of the hTR gene promoter allows the analysis of therapeutic molecules which modulate hTR promoter activity. Sequences AAZ07623-80 represents PCR primers for amplifying
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                                                                                                                                                                                                                                                                                                  Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                        Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; gene therapy; thymidine kinase gene; anticancer therapy; human;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity 100.0%; Pred. No. 78, Conservative 0; Mismatches 0; Indels
Human telomerase RNA gene (hTR) specific primer TRC3F.
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                                                                                                                                                                                                                      (CANC-) CANCER RES CAMPAIGN TECHNOLOGY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   promoter sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        46 CTAACCCTAACTGAGAAGGGCGTA 69
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                                                                                                                                                                                                                                                                                                                                        Disclosure; Fig 6; 109pp; English
                                                                                                                                                                                             98GB-00001902
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  human TR gene (hTR)
                                                    PCR primer; 88
                                                                                         Homo sapiens
                                                                                                                WO9938964-A2
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                                                                                                                                                                                                                                                                                                                 gene therapy
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                                                                                                                                         05-AUG-1999
                                                                            Synthetic
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                                                                                                                                                                                                                                                Keith WN;
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Best Local S
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The present sequence represents a peptide nucleic acid molecule which hybridises to the mRNA component of mammalian telomerase, and inhibits telomerase activity. Telomerase is a ribonucleoprotein enzyme that cynthesizes one strand of the telomeric DNA, using as a template an 11 nucleotide sequence contained within the RNA component of the enzyme. The invention relates to RNA molecules having a sequence of no more than 25 bases, which include the sequence of TARGG. The uncharged nature of the PNA ci noreases the melting temperature of associating strands, affords greater resistance of degradation by proteases or nucleases. The therapeutic PNAs may be used for treating disease conditions such as cancers, neoplasia, hyperplasia, neurodegenerative diseases. The concers, neoplasia, hyperplasia, neurodegenerative diseases, aging, human ci mamunodeficiency virus (HIV) infection/AIDS (acquired immunodeficiency syndrome) and associated pathologies, fungal infections, and other cytotoxic or diseases characterized by abnormal telomere metabolism or telomerase crivity, in combination with antineoplastic and other rototoxic or cytostatic agents, antifungal agentes, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization cytostatic agents, antifungal agentes, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization of individuals, e.g. paternity testing, based on hTR gene restriction of indoviduals, of a mammalian relomerace and a second to probes to detect the RNA comonent of a mammalian relomerace and as 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     activity
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                                                                                                                                                     /*tag= a /*tag= a /*tag= a /*tag= nucleic acid molecule, where N-(2-aminoethyl)glycine units are linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New peptide nucleic acid (PNA) compounds that inhibit telomerase activit in mammalian cells is useful as probes to detect the RNA component of a mammalian telomerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            based on pathological characteristics. The diagnostic and prognostic methods of the present invention can be used to detect an immortal or neoplastic cell or tumour tissue or cancer of any origin, provided the cell expresses telomerase activity and its RNA component
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gapв
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Shay JW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        64
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                                                                               Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (TEXA ) UNIV TEXAS SYSTEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2000-292432/25.
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les 24; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-APR-1997;
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                                                                                        Key
misc feature
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Synthetic.
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RESULT 92
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, on neurodegenerative diseases.
                                                                                                                                                                                  neurodegenerative disease, neoplasia, HIV; AIDS; cytostatic, human immunodeficiency virus, acquired immunodeficiency syndrome;
                                                                                                              Oligonucleotide #1 used in melting temperature studies of PNAs.
                                                                                                                                                                                                              telomere metabolism; anti-inflammatory; immunosuppressive; ss.
                                                                                                                                                          hTR gene RFLP pattern; cancer; inflammation; forensic;
lymphoproliferative disease; autoimmune disease; hyperplasia;
                                                                                                                                         paternity testing; human telomerase RNA component;
                                                                                                                                                                                                                                                                                                                                                                                                                                                Norton JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 24; DB 1; Length 24;
Pred. No. 78;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seguence 24 BP; 7 A; 5 C; 5 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                Corey DR,
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100.0%; Pred. No. 'c.
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                            AAS15445 standard; DNA; 24 BP.
                                                                                                                                                                                                                                                                                                                                                                          96US-00630019.
97US-00838545.
                                                                                                                                                                                                                                                                                                                                             99US-00349532
                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-638024/73
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                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                      US6294650-B1
                                                                                                                                                                                                                                                                                                                                             08-JUL-1999;
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09-APR-1997;
                                                                                  14-FEB-2002
                                                                                                                                                                                                                                                                                                                  25-SEP-2001
                                                                                                                                           Mammalian;
                                                                                                                                                                                                                                                            Synthetic.
                                                       AAS15445;
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RESULT 91
           AAS15445
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Provided are an Adenovirus vector with a telomerase promoter associated with cancer occurrence and a method for using the same for anti-cancer treatment. An Adenovirus expression vector pS-HTERT-TK is produced by using an expression cassette consisting of suicide gene HSV-TK, as late bylyadenylation signal for human telomerase hTERT promoter and bovine growth hormone BGH). Adenovirus clone Ad-hT-TK(KCCM 1038) is used for anti-cancer gene therapy. The present sequence represents a primer of the
                                                                                                                                                                                Adenovirus vector; cancer; anti-cancer; pS-hTERT-TK; suicide gene; HSV-TK; human telomerase; hTERT promoter; bovine growth hormone; BGH; Adenovirus clone; Ad-hT-TK; Cytostatic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New adenovirus vector comprising telomerase promoter, useful in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Telomerase, body fluid; cancer; tumour; screening; TRAP; diagnosis; telomeric repeat amplification protocol; detection; PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 24 BP; 7 A; 5 C; 5 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human telomerase RNA template PCR primer R3C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; SEQ ID NO 3; 1pp; Korean.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             41 TTTGTCTAACCCTAACTGAGAAGG 64
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ADQ36831 standard; DNA; 24 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                     27-JUN-2002; 2002KR-00038076.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27-JUN-2002; 2002KR-00038076.
                                                                                                                                         Primer of the invention #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAZ08704 standard; DNA; 25
                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    treatment of cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SONG J S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 KIM Y T.
                                                                                                                                                                                                                                                                                                                            KR2004002322-A.
                                                                                                                                                                                                                                                                              Unidentified,
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                                                                                           26-AUG-2004
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                                                ADQ36831;
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64

TTTGTCTAACCCTAACTGAGAAGG 24

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Wright
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                                                                                                                                                                                 cellular portion of body fluid from a cancer patient using a modified cellular portion of body fluid from a cancer patient using a modified tellular portion of body fluid from a cancer patient using a modified tellular proteocal (TRAP). A method for detecting cancer comprises: (a) removing the cellular portion of a body fluid specimen from the patient; (b) preparing a protein extract from the body fluid remainder; (c) assaying the extract for the presence and quantity of tellomerase RNA or telomerase activity; and (d) comparing the results with normal levels, to determine the presence of cancer. The methods are used in cancer diagnosis and prognosis, and also to monitor cancer the rappy effectiveness. Unlike prior art telomerase activity assays in cancer patients, the method allows noninvasive sample collection. The methods are also more reliable and less tumour specific than other methods which detect circulating tumour markers. The present sequence represents a human telomerase RNA template PCR primer used in the exemplification of the present invention
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                                                                                                                                Detecting telomerase activity in non-cellular body fluid using a modified telomeric repeat amplification protocol.
                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Peptide nucleic acid; PNA; cancer; telomerase; probe; hybridisation; inhibitor; human telomerase RNA; hTR; PCR; oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA oligonucleotide 2, used in the measurement of Tm values.
                                                                                                                                                                                                                                                                                                                                                                          Score 24; DB 1; Length 25;
Pred. No. 82;
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                                                                                          Abruzzo LV;
                                                                                                                                                                                                                                                                                                                                                        Sequence 25 BP; 7 A; 3 C; 9 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                     5.3%; Scor.
100.0%; Pred. No. c...
0; Mismatches
                                                                                          Highsmith E,
                                                                                                                                                                                                                                                                                                                                                                                                                      CTTCCACCGTTCATTCTAGAGCAA 168
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                                                                                                                                                                  Disclosure; Page 16; 32pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                 CTTCCACCGTTCATTCTAGAGCAA
                                                                     (UYMA-) UNIV MARYLAND BALTIMORE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP
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                              99WO-US003302
                                                  98US-0074793P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                          Stamberg J,
                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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                                                                                                                WPI; 1999-508655/42.
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                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                              16-FEB-1999;
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          19-AUG-1999
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                                                                                                                                                 their
used
                                                                                                                                            This is an oligonucleotide used in the measurement of Tm values and thei complimentary peptide nucleic acids (PNAs), (e.g. AAT89225-T89227). PNAs hybridise specifically to an RNA component of mammalian telomerase, and include the sequence GGG for specific hybridisation to the template region of this component. PNAs can be used as probes to detect the RNA component of mammalian telomerase and as inhibitors of telomerase activity, especially in the treatment of cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "Peptide nucleic acid molecule, where N-(2-aminoethyl)glycine units are linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence represents a peptide nucleic acid molecule which hybridises to the mRNA component of mammalian telomerase, and inhibits telomerase activity. Telomerase is a ribonucleoprotein enzyme that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Peptide nucleic acid; PNA; telomerase; ribonucleoprotein enzyme; candinhibitor; neoplasia; neurodegenerative disease; aging; hyperplasia; AIDS; HIV; fungal infection; forensic identification; detect; tumour; paternity testing; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
nucleic acids hybridising to mammalian telomerase RNA -
                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DR;
                                                                                                                                                                                                                                                                                                                                                                                                      DB 1; Length 23;
89;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indele
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Corey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PNA sequence #26 used to inhibit telomerase activity.
                                                                                                                                                                                                                                                                                                                                                        Sequence 23 BP; 5 A; 7 C; 1 G; 10 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Norton JC,
                                                                                                                                                                                                                                                                                                                                                                                                   5.1%; Score 23; DB
100.0%; Pred. No. 89;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Shay JW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                  Example 2; Page 49; 76pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 2; Col 33; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ВP
                                                  diseases, also for diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              97US-00838545.
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                       telomerase, for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAA37568 standard; DNA; 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Piatyszek MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (TEXA ) UNIV TEXAS SYSTEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                        23; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mammalian telomerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2000-292432/25
                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 39-APR-1997;
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New peptide
to inhibit t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
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conclocide sequence contained within the RNA component of the enzyme. The invention relates to PNA molecules having a sequence of no more than 25 bases, which include the sequence GTTAGG. The uncharged nature of the PNA bases, which include the sequence GTTAGG. The uncharged nature of the PNA backbone increases the rate of association with targeted nucleic acids, and affords greater resistance of degradation by proteases or nucleases. The increases the rate of degradation by proteases or nucleases. The cherapeutic PNAs may be used for treating diseases or nucleases. The immunodeficiency virus (HIV) infection/AIDS (acquired immunodeficiency consection virus and other rucleocides why be cateforded for molecular diagnostics, labelled MNAs are used as whysiciation or individuals, e.g. paternity testing, based on hTR gene restriction fragment length polymorphism (RRLP) pattern. PNAs are also useful as probes to detect the RNA component of a mammalian telomerase and as probes to detect the RNA component of a mammalian telomerase and as concerous conditions to be detected with increased confidence and allows cancerous conditions to be detected with increased confidence and conditions to be detected with increased conf
synthesizes one strand of the telomeric DNA, using as a template an 11\, nucleotide sequence contained within the RNA component of the enzyme.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 23 BP; 5 A; 7 C; 1 G; 10 T; 0 U; 0 Other;
                $$$$$$$$$$$$$$$$$$$$$$$$$$$$$$$$
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ö Gaps 0 Score 23; DB 1; Length 23; Pred. No. 89; 0; Indels Mismatches 35 CCATTTTTGTCTAACCTAACT 57 1 CCATTTTTGTCTAACCCTAACT 23 / Match 5.1%; Sc. Local Similarity 100.0%; Pi nes 23; Conservative 0; Query Match Best Loca Matches 셤

AAS15446 standard; DNA; 23 (first entry) 14-FEB-2002 AAS15446; RESULT 96 AAS15446

hTR gene RPLP pattern; cancer; inflammation; forensic; lymphoproliferative disease; autoimmune disease; hyperplasia; lymphoproliferative disease; neoplasia; HIV; AIDS; cytostatic; human immunodeficiency virus; acquired immunodeficiency syndrome; telomere metabolism; anti-inflammatory; immunosuppressive; ss. Oligonucleotide #2 used in melting temperature studies of PNAs. Mammalian; paternity testing; human telomerase RNA component;

99US-00349532. 96US-00630019. 97US-00838545. sapiens. US6294650-B1. 08-JUL-1999; 09-APR-1996; 09-APR-1997; 25-SEP-2001 Synthetic. Homo

Norton JC; Shay JW, Wright WE, Piatyszek MA, Corey DR, WPI; 2001-638024/73

(TEXA) UNIV TEXAS SYSTEM.

ö The present invention relates to peptide nucleic acids (FNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in mammalian cells by hybridising to the RNA component of mammalian telomerase. The PNAs are useful as probes to detect the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polynucleotide having the human telomerase RNA component (hTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the hTR gene RFLP pattern. The suspects or unknown descendants based on the hTR gene RFLP pattern. The Cytostatic; vasotropic; protozoacide; immunosuppressive, dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; double-stranded RNA; micro-RNA; miRNA; short halfipin RNA; shrNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss. PNA can be further used for treating or preventing cancer, inflammation. Iymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmaceuticals (such as antineoplastic or cytostatic agents) can be used for treating neoplasia, hyperplasia, human immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomere metabolism or telomerase activity. The present sequence representing a DNA oligonucleotide is complementary to some of the PNAs of the present invention, and is used Gaps New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. .; DB 1; Length 23; 89; 0; Indels Human TERC mRNA transcript target sequence, SEQ ID 521. Sequence 23 BP; 5 A; 7 C; 1 G; 10 T; 0 U; 0 Other; Local Similarity 100.0%; Pred. No. 89; es 23; Conservative 0; Mismatches Score 23; 57 1 ccarrrrrgrcraacccraacr 23 Example 2; Col 34; 46pp; English. 35 CCATTTTTGTCTAACCCTAACT n melting temperature studies ADF93794 standard; mRNA; 23 BP 2002US-0396600P. 2002US-0406784P. 2002US-0408378P. 2002US-0409293P. 2002US-0363124P. 2002US-0386782P. 11-FEB-2003; 2003WO-US004088 2002US-0358580P 5.1%; 26-FEB-2004 (first entry) WO2003070742-A1. 29-AUG-2002; 05-SEP-2002; 09-SEP-2002; Homo sapiens. 20-FEB-2002; 11-MAR-2002; 06-JUN-2002; 17-JUL-2002; 28-AUG-2003 ADF93794; Query Match Matches RESULT 97 ADF93794 ò g

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                                                                                                                                                                                                                   The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, expressed from a vector or enzymatically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA, and vectors that express siNA. The siNAs are
                                                                                                                                                                                                                                                                                                                                                                                                                                used to modulate expression of the telemerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNhs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short halfpin RNA; shRNA; expression medulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                  New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human TERT mRNA transcript target sequence, SEQ ID 530.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             mapping (e.g., of single nucleotide polymorphisms). represents a human TERC transcript target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 23 BP; 6 A; 7 C; 3 G; 0 T; 7 U; 0 Other;
                                                                                                                                                                                        Example 3; SEQ ID NO 521; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      146 TTCCACCGTTCATTCTAGAGCAA 168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADF93803 standard; mRNA; 23 BP
15-JAN-2003; 2003US-0440129P
                                (RIBO-) RIBOZYME PHARM INC
                                                                  Mcswiggen J, Beigelman L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-FEB-2004 (first entry)
                                                                                                   WPI; 2003-689777/65.
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The invention relates to short interfering nucleic acids (sink) which downregulate expression of the one or more telomerase genes by RNA interference. The sinks may or may not comprise sense and antisense of adouble or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the sinks include short interfering RNA (siRNA), double stranded RNA, micro-RNA (miRNA) and short can contain deoxyribonucleotides, and can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, can complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treatment rejection, or autoimmune or age-related diseases, cortex marging datasensis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening datagnosis, lupus erythematosus, AIDS, macular degeneration, canceric engineering, pharmacogenomics, studying gene function, and gene capersents a human TERT transcript target sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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Pred. No. 89;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 23 BP; 6 A; 10 C; 6 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure, SEQ ID NO 530, 145pp, English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5.1%; Score 23;
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                                                                                               11-MAR-2002; 2002US-0363124P.

06-UTV-2002; 2002US-03867B2P.

17-UTV-2002; 2002US-0396600P.

29-AUG-2002; 2002US-0406784P.

05-SEP-2002; 2002US-040837BP.

05-SEP-2002; 2002US-040933P.

15-JAN-2003; 2003US-0440129P.
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11-FEB-2003; 2003WO-US004088
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                                                                                                                                                                                                                                                                                                                                                                                                    (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mcswiggen J, Beigelman L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-FEB-2004 (first entry)
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Matches 22, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-689777/65.
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonuclectide and an antisense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised, can contain deoxyribonuclectides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and teaded for treating cancer, restencis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, can be entily, diagnosis, therapeutic target identification and validation, contain and validation, contain and validation, contain and relating, diagnosis, therapeutic target identification and validation, contains and relating or entily.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence
drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mapping (e.g., of single nucleotide polymorphisms). represents a human TERC transcript target sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 3; SEQ ID NO 522; 145pp; English.
                                                                                                                                                                                                                                                                                  2002US-0386782P.
2002US-0396600P.
2002US-0406784P.
2002US-0408378P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                           (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Beigelman L;
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05-SEP-2002;
09-SEP-2002;
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06-JUN-2002;
17-JUL-2002;
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                                                                      Homo sapiens
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The invention relates to short interfering nucleic acids (siNA) which compures developments of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense corporates or alternatively are assembled from a sense onigonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double stranded RNA, micro-RNA (miRNA) and short can contain deoxyribonucleotides, and can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised.

CC expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue cappants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically conditions) and unlipple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, lupus erythematosus, AIDS, macular degeneration, cappaning, diagnosis, lupus erythematosus, AIDS, macular degeneration, cappaning (e.g., of single nucleotide polymorphisms). The present sequence mapping (e.g., of single nucleotide polymorphisms). The present sequence
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Pred. No. 89;
7; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 69.6%;
Matches 16; Conservative
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5; Mismatches
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RESULT 100

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ADF93793

SXXXE

Matches

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Gaps

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23

DB 1; Length 23; 0; Indels

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Cytostatic, vasotropic, protozoacide, immunosuppressive, dermatological; neuroprotective, anti-HIV; ophthalmological; antiulcer; anti-rheumatic;
                                                                                                                         antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; mitch are short interfering RNA; siRNA; double-stranded RNA; mitch miRNA; mirch hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
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Human TERC mRNA transcript target sequence, SEQ ID 520.
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2002US-0396600P.
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05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409239P.
15-UAN-2003; 2003US-0440129P.
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06-JUN-2002;
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US09436060A-16.rng.sl

Sequence 23 BP; 2 A; 11 C; 4 G; 0 T; 6 U; 0 Other;

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51

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can complexes of siNA, and vectors that express siNA. The siNAs are and/or complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue caplants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically proteosal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematoguus, AlbS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                           Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antimheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; slNA; short interfering RNA; siRNN; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
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                                                                                      ADF93802 standard; mRNA; 23 BP.
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2002US-0396600P.
2002US-0406784P.
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2002US-0409293P.
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                                                                                                                                                                         ADF93802;
RESULT 101
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The invention relates to short interfering nucleic acids (sinA) which downregulate expression of the one or more telomerase genes by RNA interference. The sinAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antidense regions, or alternatively are assembled from a sense oligonuclectide and an integense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised. Cexpressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are and/or complexes of siNA; and vectors that express siNA. The siNAs are cused to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be
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                                              Gaps
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        Score 23; DB 1; Length 23; Pred. No. 89;
                                              Indels
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                                              6; Mismatches
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        Score 23;
                                                                                   136 GCCTGCCGCCTTCCACCGTTCAT 158
                                                                                                         1 GCCUGCCGCCUUCCACCGUUCAU 23
                                                                                                                                                                                                                       ADF93804 standard; mRNA; 23 BP.
    5.1%;
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2002US-0363124P.
2002US-0386782P.
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Ouery Match
Best Local Similarity 73.9°
Matches 17; Conservative
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                 protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siMAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents a human TERT transcript target sequence.
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   treating cancer, restenosis, infectious diseases (specifically
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Pred. No.
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82.6%;
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2002US-0408378P.
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2002US-0386782P.
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                                                                                                                                                                                               Query Match
Best Local Similarity 82.6<sup>5</sup>
Matches 19, Conservative
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29-AUG-2002;
05-SEP-2002;
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   used for
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                                                                                                                                                                                                                                                                                                                                                                        RESULT 103
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interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised.

Expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNas are used to modulate expression of the telomerase genes in cells, tissue captains or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restencisis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, comultiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering pharmacogenomics, studying gene function, and gene mapping (e.g., of single mucleotide polymorphisms). The present sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or activity.
                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           double-stranded short interfering nucleic acid; siNA; antibarkinsonian; antiarteriosclerotic; neuroprotective; nootropic; antiparkinsonian; anticonvulsant; pulmonary disease; restenosis; atherosclerosis; Alzheimer's; Parkinson's; epilepsy; dementia; huntington's; amyotrophic lateral sclerosis; gene therapy; target; ss; hTR.
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                                                                                                                                                                                                                                                                                                                                                             DB 1; Length 23;
                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                        Sequence 23 BP; 2 A; 3 C; 14 G; 0 T; 4 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                              4; Mismatches
                                                                                                                                                                                                                                                                                                                                                         Score 23; DB
Pred. No. 89;
                                                                                                                                                                                                                                                                                                                                                                                                                                2 GGTTGCGGAGGGTGGGCCTGGGA 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               hTR siNA-target RNA - SEQ ID 92.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (SIRN-) SIRNA THERAPEUTICS INC
                                                                                                                                                                                                                                                                                                                                                             5.1%;
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2002US-0409293P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                            19; Conservative
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                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
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29-AUG-2002;
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Example 24; SEQ ID NO 85; 593pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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                                                                The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 1 nucleotides. The siNA of the invention demonstrates antiarteriosclarctic. The siNA of nootropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary diseases such as restenosis, atherosclerosis, Alzheimer's disease, Parkinson's disease, epilepsy, dementia, huntington's disease or amyotrophic lateral sclerosis. Purthermore, the siNA may be utilised for penerapy applications. The current sequence is that of the siNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        double-stranded short interfering nucleic acid; siNA;
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Example 24; SEQ ID NO 92; 593pp; English.
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2002US-0363124P.
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2002US-0409293P.
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11-MAR-2002;
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Matches
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                               (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of
The invention relates to a double-stranded short interfering nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                  for
                                                                                                                               the invention demonstrates antiarteriosclerotic, neuroprotective, nootropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian tharget gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary diseases such as restenosis, atherosclerosis, Alzheimer's disease, Parkinson's disease, epilepsy, dementia, huntington's disease or amyotrophic lateral sclerosis. Furthermore, the siNA may be utilised fogene therapy applications. The current sequence is that of the siNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1; Length 23;
89;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5.1%; Score 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pred. No
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 GGTTGCGGAGGGTGGGCCTGGGA 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Thompson J;
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2002US-0408378P.
2002US-0409293P.
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2002US-0363124P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              target DNA of the invention.
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Matches 19; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-731676/69.
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The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of
Example 24; SEQ ID NO 87; 593pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Beigelman L,
                                                                                                                                                                                                                                                                                                                                                                                                           ADG29524 standard; RNA; 23
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                                                                                                                                                                                                                                                                                       Matches 22; Conservative
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Gran N, Teman N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-731676/69.
                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
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05-SEP-2002;
09-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unidentified
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06-JUN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                 RESULT 108
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                         The invention relates to a double-stranded short interfering nucleic acid (sink) molocule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of
                                                                         the invention demonstrates nutiatreriosclerotic, neuroprotective, nootropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary diseases such as restenoses, atherosclerosis, Albrahimer's disease, Parkinson's disease, epilepsy, dementia, huntington's disease or amyotrophic lateral sclerosis. Furthermore, the siNA may be utilised for gene therapy applications. The current sequence is that of the siNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or activity.
                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             double-stranded short interfering nucleic acid; siNA; antiarteriosclerotic; neuroprotective; nootropic; antiparkinsonian; anticonvulsant; pulmonary disease; restenosis; atherosclerosis; Alzheimer's; Parkinson's; epilepsy; dementia; huntington's; amyotrophic lateral sclerosis; gene therapy; target; ss; hTR.
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                                                                                                                                                                                                                                                            Score 23; DB 1; Length 23;
Pred. No. 89;
7; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pavco P,
                                                                                                                                                                                                                                    Sequence 23 BP; 6 A; 7 C; 3 G; 0 T; 7 U; 0 Other;
  Example 24; SEQ ID NO 91; 593pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chowrira
                                                                                                                                                                                                                                                                                                                  146 TTCCACCGTTCATTCTAGAGCAA 168
                                                                                                                                                                                                                                                                                                                                Thompson J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hTR siNA-target RNA - SEQ ID 87.
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2002US-0386782P.
2002US-0406784P.
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                                                                                                                                                                                                           target DNA of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                            ADG29521 standard; RNA; 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                            Local Similarity 69.6
nes 16; Conservative
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Q. Usman N,
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06-JUN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unidentified
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Jamison S,
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                                                                                                                                                                                                                                                              Query Match
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the invention demonstrates antiarteriosclerotic, neuroprotective, nootropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regularing the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary diseases such as restenosis, atherosclerosis, Albreimer's disease, Parkinson's disease, epilepsy, dementia, huntington's disease or amyotrophic lateral sclerosis. Butthermore, the siNA may be utilised for gene therapy applications. The current sequence is that of the siNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antiarteriosclerotic; neuroprotective; nootropic; antiparkinsonian; anticonvulsant; pulmonary disease; restenosis; atherosclerosis; Alzheimer's; Parkinson's; epilepsy; dementia; huntington's; amyotrophic lateral sclerosis; gene therapy; target; ss; hTR.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1; Length 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            double-stranded short interfering nucleic acid; siNA;
                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 23 BP; 6 A; 10 C; 6 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             89;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 23;
Pred. No. 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     283 GCACCCACTGCCACCGCGAAGAG 305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 GCACCCACUGCCACCGCGAAGAG 23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       hTR siNA-target RNA - SEQ ID 90.
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2002US-0408378P.
2002US-0409293P.
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2002US-0363124P.
2002US-0386782P.
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                                                                                                                                                                                                                                                                                                                                                        target DNA of the invention.
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Example 24; SEQ ID NO 86; 593pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                            ADG29522;
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of the
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                   The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of the invention demonstrates antiarteriosclerotic, neuroprotective, nootropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition way include pulmonary diseases such as restenosis, atherosclerosis, Alzheimer's disease, parkinson's disease, epilepsy, dementia, huntington's disease, publications. Furthermore, the siNA may be utilised for demonstrappy applications. The current sequence is that of the siNA
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                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antiarteriosclerotic, neuroprotective, nootropic; antiparkinsonian, anticonvulsant; pulmonary disease; restenosis; atherosclerosis; Alzheimer's; Parkinson's; epilepsy; dementia; huntington's; amyotrophic lateral sclerosis; gene therapy; target; ss; hTR.
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                                                                                                                                                                                                                                     5.1%; Score 23; DB 1; Length 23; 69.6%; Pred. No. 89; 1; Mismatches 0; Indele
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 double-stranded short interfering nucleic acid; siNA;
                                                                                                                                                                                                             Sequence 23 BP; 4 A; 9 C; 3 G; 0 T; 7 U; 0 Other;
Example 24; SEQ ID NO 90; 593pp; English.
                                                                                                                                                                                                                                                                                     144 CCTTCCACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                 1 CCUUCCACCGUUCAUUCUAGAGC 23
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2002US-0408378P.
2002US-0409293P.
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2002US-0363124P.
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                                                                                                                                                                                       target DNA of the invention.
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                                                                                                                                                                                                                                                            16; Conservative
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                                                                                                                                                                                                                                                 Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unidentified
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                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                            RESULT 109
ADG29520
                                                                                                                                                                                                                                                             Matches
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                                (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of
The invention relates to a double-stranded short interfering nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                 for
                                                                                                                       the invention demonstrates antiarteriosclerotic, neuroprotective, neotropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, rissue or organism. The disease or condition may include pulmonary diseases such as restenosis, atherosclerosis, Alzheimer's disease or amyorophic lateral sollegy, dementia, huntington's disease or amyorophic lateral sollegis. Furthermore, the sink may be utilised for gene therapy applications. The current sequence is that of the sink target DNA of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1; Length 23;
89;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 23 BP; 2 A; 11 C; 4 G; 0 T; 6 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5.1%; Score 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred. No
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      136 GCCTGCCGCCTTCCACCGTTCAT 158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GCCUGCCGCCUUCCACCGUUCAU 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Thompson J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TR SINA-target RNA - SEQ ID 88.
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2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
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2002US-0363124P.
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nes 17; Conservative
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gene therapy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                 AAZ07300;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Keith WN;
                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                  RESULT 112
                                                                                                                                                                                                                          Matches
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                  The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of the invention demonstrates antiatrefricoslerotic, neuroprotective, cusful for down-regulating the expression of an endogenous mammalian that responds to modulation of gene expression of any disease or condition that responds to modulation of gene expression or activity in a cell, cissue or organism. The disease or condition may include pulmonary diseases such as restenosis, atherosclerosis, Alzheimer's disease, pilepsy, dementia, huntington's disease or comparations of gene therepy applications. The current sequence is that of the sina
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention relates to a novel method concerning prognostic markers associated with EGFR (epidermal growth factor receptor) positive cancer. Specifically, it refers to a gene expression profiling method that can provide a prediction as to whether a patient is likely to respond well to treatment with an EGFR inhibitor. The present invention describes the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Prognosing a patient with EGFR-expressing colon cancer comprises subjecting a sample comprising EGFR-expressing cancer cells to quantitative analysis of the expression level of the RNA transcript of at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR primer to amplify a human cancer prognostic marker DNA SeqID 345.
                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           human; primer; PCR; prognostic marker; EGFR; epidermal growth factor receptor; cancer; gene expression profiling; microarray; head and neck cancer; colon cancer; metastatic spread;
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                                                                                                                                                                                                                           Sequence 23 BP; 2 A; 8 C; 9 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                Pred. No. 89;
4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Shak S, Baselga J;
 Example 24; SEQ ID NO 88; 593pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 54; SEQ ID NO 345; 113pp; English.
                                                                                                                                                                                                                                                                                                      395 GCGCGCGCGATTCCCTGAGCTG 417
                                                                                                                                                                                                                                                                                                                     GCGCGGCGCAUUCCCUGAGCUG 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (VALL-) VALL HEBRON UNIV HOSPITAL
                                                                                                                                                                                                                                                                                                                                                                                             ADP27908 standard; DNA; 23 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14-NOV-2003; 2003WO-US036777.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-NOV-2002; 2002US-0427090P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  least one gene e.g., CD44v3.
                                                                                                                                                                                                    target DNA of the invention.
                                                                                                                                                                                                                                                                  82.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (GENO-) GENOMIC HEALTH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   neoplastic disease; ss
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2004-420643/39.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                              26-AUG-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                     ADP27908;
                                                                                                                                                                                                                                                                                Matches
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quantitative analysis of the expression level of the RNA transcript of at least one gene selected from the group of CD44v3, CD44v6, DR5, GR01, KRT17, LAMC2 or their products thereof. It further provides a CDNA microarray containing named genes that represent prognostic transcripts which are useful for determining whether a patient diagnosed with an EGFR expressing head or neck cancer or colon cancer shibits elevated or decreased expression levels of these genes compared to normal. As such, these methods are also useful for prognosing or predicting the likelihood of cancer-attributable death or progression, including recurrence and metasteric spread of a neoplastic disease, as well as drug resistence. This oligonucleotide sequence is a PCR primer used to amplify a human PCR amplicon DNA sequence used as a prognostic cancer marker, given in an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; gene therapy; thymidine kinase gene; anticancer therapy; human; mutagenesis; PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1; Length 23;
89;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 23 BP; 6 A; 9 C; 5 G; 3 T; 0 U; 0 Other;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       404 GATTCCCTGAGCTGTGGGACGTG 426
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23 GATTCCCTGAGCTGTGGGACGTG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 exemplification of the invention.
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100.0%;
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Les 23; Conservative
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Sequence 27 BP; 2 A; 4 C; 12 G; 9 T; 0 U; 0 Other;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RNA components of (non)human mammalian telomerase(s) - useful in studying cell senescence and immortalisation.
(especially pGT62-codAupp) comprising the promoter and a viral thymidine kinase gene can be used to convert a prodrug, e.g. gancyclovir, so that neoplasia can be controlled or treated. Direct down-regulation of telomerase RNA gene through manipulation of transcription factors may be effective anticancer therapy and the cloning of the hTR gene promoter allows the analysis of therapeutic molecules which modulate hTR promoter activity. Sequences AAZO7696-321 represent PCR primers used in cloning and mutagenisis of human TR gene (hTR) promoter region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The RNA components of (non) human mammalian telomerase(s) especially from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Telomerase; mammal; antisense; triplex forming oligonucleotide; plasmid;
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100.0%; Pred. No. 1e+02;
iive 0; Mismatches 0; Indels
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                                                                                                                                                          Sequence 25 BP; 1 A; 5 C; 15 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Primer used for producing telomerase probe.
                                                                                                                                                                                                                                                                                  GGGTTGCGGAGGGTGGGCCTGGG 25
                                                                                                                                                                                                                                                             1 GGGTTGCGGAGGGTGGGCCTGGG 23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SPRING HARBOR LAB
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95US-00387524.
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                                                                                                                                                                                                            Best Local Similarity
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GERON CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9601614-A2
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13-FEB-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         - useful in studying
                                                                                                                                                                                                                                                                                                      Telomerase; mammal; antisense; triplex forming oligonucleotide; plasmid;
                                Gaps
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                                                                                                                                                                                                                                                                       Antisense oligonucleotide (P3) inhibiting telomerase activity.
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 Length 27;
                                Indels
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Villeponteau B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RNA components of (non)human mammalian telomerase(s)
DB 1; Ler
1.1e+02;
hes 0;
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Query Match 5.1%; Score 23; DB Best Local Similarity 100.0%; Pred. No. 1.1 Matches 23; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cell senescence and immortalisation.
                                                                44
                                                                                            GGAGGGGTGGTGCCATTTTTG 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 23; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          46 CTAACCCTAACTGAGAAGGGCG 67
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                                                                22 GGAGGGGTGGTGGCCATTTTTG
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95US-00387524.
95US-00485778.
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                                                                                                                                                                           BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                94US-00272102
                                                                                                                                                                                                                                                                                                                        probe; primer; ribozyme; ss.
                                                                                                                                                                           AAT11033 standard; DNA; 22
                                                                                                                                                                                                                                         02-JUL-1996 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1996-097428/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          * Best Local Similarity Matches 22; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GERON CORP
                                                                                                                                                                                                                                                                                                                                                                                    WO9601614-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-OCT-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13-FEB-1995;
07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                   25-JAN-1996
                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                          AAT11033;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             * "Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (COLD-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GERO-)
                                                                                                                                           RESULT 114
                                                                                                                                                          AAT11033/
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                                                                                            유
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RNA component of mammalian telomerase, esp. human - useful in identifying
                                                                                                                                                                                                                                                                                                                                                            The present sequence is a RNA component of mammalian telomerase, antisense oligonucleotide, which can be used, along with triple helix forming sequences, to inhibit telomerase activity in cells, esp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RNA component of mammalian telomerase antisense oligonucleotide P3.
         RNA component; mammalian; telomerase; antisense oligonucleotide;
triple helix; inhibition; neoplastic; cells; activity; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RNA component; mammalian; telomerase; antisense oligonucleotide;
triple helix; inhibition; neoplastic; cells; activity; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 22; DB 1; Length 22;
Pred. No. 1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 22 BP; 2 A; 9 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Villeponteau B, Feng J, Funk W, Andrews WH;
                                                                                                                                                                                                                                             Andrews WH;
                                                                                                                                                                                                                                                                                                             e.g. candidate telomerase-modulating agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 38; 114pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          75
                                                                                                                                                                                                                                             Funk W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22 AACTGAGAAGGCGGTAGGCGCC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                      4.9%; Scallarity 100.0%; Pi
Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          54 AACTGAGAAGGCGTAGGCGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 94US-00272102.
94US-00330123.
95US-00472802.
95US-00482115.
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94US-00330123.
95US-00472802.
95US-00482115.
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                                                                                                                                                                                                                                             Villeponteau B, Feng J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAT10287 standard; DNA;
                                                                                                                                                                                                                                                                       WPI; 1996-097581/10.
                                                                                                                                                                                                                    (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
Matches 22; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                      neoplastic cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GERO- ) GERON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9601835-A1.
                                                                                                                            06-JUL-1995;
                                                                          WO9601835-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-JUL-1995;
                                                                                                                                                                 27-OCT-1994;
07-JUN-1995;
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                                                                                                                                                                                            07-JUN-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-JUN-1995
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                                                                                                    25-JAN-1996
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                                                  Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The RNA components of (non) human mammalian telomerase(s) especially from mouse, rat and chinese hamster are all claimed. Antisense obligonucleotides can be used to block the activity of the telomerase; probes and primers can be used in detection; vectors and host cells transformed with the isolated telomerase genes can be used for production of telomerases; RNA and DNA ribozymes and triplex forming oligonucleotides directed against the telomerase genes can be used therapeutically as can plasmids. A mouse which lacks the telomerase gene therapeutically as can plasmids. A mouse which lacks the telomerase gene the role it plays in immortalisation. The antisense oligonucleotide is synthesised as a 2-0-methyl RNA oligonucleotide and is more resistant to hydrolysis than unmodified RNA oligonucleotides (See AATI1032-35)
                                                                                                                                                                                                                                                                                                                                                                                                                                                          components of (non)human mammalian telomerase(s) - useful in studying
                                                                                                                                          Telomerase; mammal; antisense; triplex forming oligonucleotide; plasmid;
                                                                                                                                                                                                                                                                                                                                                                                          Marhuenda MA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RNA component of mammalian telomerase antisense oligonucleotide TA3.
                                                                                                                Antisense oligonucleotide (TA3) inhibiting telomerase activity.
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0
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100.0%; Pred. No. 1e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                          Greider C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 22 BP; 2 A; 9 C; 5 G; 0 T; 6 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                          Funk W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cell senescence and immortalisation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 23; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         54 AACTGAGAAGGGCGTAGGCGCC 75
                                                                                                                                                                                                                                                                                                                                                                                           Feng J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AACTGAGAAGGCGTAGGCGCC 1
                                                                                                                                                                                                                                                                                   94US-00272102.
94US-00330123.
95US-00387524.
95US-00485778.
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                                     AAT11034 standard; DNA; 22 BP.
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                                                                                                                                                                                                                                                            95WO-US008620
                                                                                                                                                       probe; primer; ribozyme; ss
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                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                  Villeponteau B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1996-097428/10.
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                                                                                                                                                                                                          WO9601614-A2
                                                                                                                                                                                                                                                            07-JUL-1995;
                                                                                                                                                                                                                                                                                                27-OCT-1994;
13-FEB-1995;
                                                                                        02-JUL-1996
                                                                                                                                                                                                                                                                                                                            07-JUN-1995;
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                                                                                                                                                                                                                                   25-JAN-1996
                                                                                                                                                                                 Synthetic.
                                                             AAT11034
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                                                                                                                                                                                                                                                                                                                                                    (COLD-)
                                                                                                                                                                                                                                                                                                                                                                 (GERO-)
          Matches
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22
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Matches
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8×8338
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                               RNA component of mammalian telomerase, esp. human - useful in identifying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RNA component of mammalian telomerase, esp. human - useful in identifying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence, a return primer for the RNA component of human telomerase (RCHT), was used in a primed in situ labelling (RRINS) procedure. The RVHT can be used in the recombinant prodm of an active telomerase mol., capable of adding sequences to chromosomal DNA telomeres, and in the synthesis of mutant sequences for the detection of mutant mammalian telomerase RNA component polynucleotides. The RCHT may also be used in the identification of telomerase modulating agents, and in the detection of telomerase related, or neoplastic conditions in a patient. Polynucleotides of at least 25 consecutive nucleotides
                                                                                    The present sequence is a RNA component of mammalian telomerase, antisense oligonucleotide, which can be used, along with triple helix forming sequences, to inhibit telomerase activity in cells, esp. neoplastic cells
                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                  recombinant production, synthesis, mutant, detection, mammalian, identification, modulating agent, neoplastic condition; transcriptional regulatory sequence, gene therapy, disease;
                                                                                                                                                                                   ö
                                                                                                                                                             4.9%; Score 22; DB 1; Length 22; 100.0%; Pred. No. 1e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                         component; human; telomerase; return primer; PRINS;
                                                                                                                                                                                                                                                                                                                                                   RNA component of human telomerase PRINS return primer.
                                                                                                                                       Sequence 22 BP; 3 A; 6 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Andrews WH;
                                          candidate telomerase-modulating agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                candidate telomerase-modulating agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 13; Page 90; 114pp; English.
                                                              Disclosure, Page 38; 114pp; English
                                                                                                                                                                                                        46 CTAACCCTAACTGAGAAGGGCG 67
                                                                                                                                                                                                                    CTAACCCTAACTGAGAAGGGCG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Funk W,
                                                                                                                                                                                                                                                                       AAT10308/c
ID AAT10308 standard; DNA; 22 BP.
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95US-00472802.
95US-00482115.
                                                                                                                                                                                                                                                                                                                                                                                                                     primed in situ labelling; ss
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                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                  22; Conservative
         WPI; 1996-097581/10.
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                                                                                                                                                                        Local Similarity
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07-JUN-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                            22
                                                                                                                                                                                                                                                                                                        AAT10308;
                                                                                                                                                              Query Match
                                           e.g.
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                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                         RNA
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The present sequence represents PCR 3'-primer hal88 used for amplifying the human telomerase (hTR). The RNA and DNA can be used in hybridisation assays to detect or quantify telomerase activity in cells, tissue or fluid samples, e.g. for diagnosis of eukaryotic parasites (yeast and protozoa) or tumours. It is also useful as primers for amplification assays. The truncated or recombinant vertebrate telomerase is used therapeutically to increase telomerase activity (also as reagants in the screening assay) while the RNA or other inhibitors such as antisense molecules, are used to reduce such activity. Typical applications are infitation/restoration of activity to cause sensescence or to prevent immortalisation of cells in tumours or parasites. The DNA is also used to raise antibodies for diagnostic detection of telomerase. Detecting telomerase allows early diagnosis of tumour or infection, before clinical slowled cause fewer side effects than drugs currently used to treat such infections. The DNA is encodes those parts of them and the contract of the parts of treat such infections. The DNA is encodes those parts of them and the creat such infections.
                                                                                                                                                                                                                                                                                                         ö
identical, or complementary to the RCHT sequence linked to heterologous transcriptional regulatory sequences, can be used for the gene therapy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cancer; eukaryotic parasite; hTR; vertebrate telomerase; yeast; protozoa; tumour; antibody; polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA encoding essential RNA components of human telomerase - also truncated or recombinant telomerase, useful for diagnosis and treatment of cancer and infection by eukaryotic parasites.
                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        out are significantly shorter than the endogenous RNA component
                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                         4.9%; Score 22; DB 1; Length 22; 100.0%; Pred. No. 1e+02; ive 0; Mismatches 0; Indels
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                                                                                                                                                        Sequence 22 BP; 5 A; 5 C; 12 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human telomerase PCR 3'-primer hal88.
                                                                                                                                                                                                                                                                                                                                                                                     183 CTGCTGGCCCGTTCGCCCCTCC 204
                                                                                                                                                                                                                                                                                                                                                                                                                                  CIGCIGGCCCGTICGCCCCTCC 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1997-099928/09
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Best Local Similarity
                                                                                                                                                                                                                                                                 Local Similarity
nes 22; Conserv
                                                                            human diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              06-JUN-1996;
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Matches

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This invention describes a novel composition comprising human telomerase having at least 2000-fold (preferably at least 6000-fold) increased relative purity compared with crude extract of cells from adenovirus transformed kidney cell line. The composition is useful for eliciting an immune response in animals and may therefore be used as a vaccine for tracting diseases associated with the overexpression of telomerase e.g. cancer. AAZ23656-Z23637 represent oligonucleotides used in the isolation of human clone 28-1 which contains a fragment of the human telomerase
                                                                                                                                                                                           Telomerase; human; immune response; cancer; vaccine; treatment; disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Compositions comprising human telomerase, useful for treating diseases associated with overexpression of telomerase e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antisense oligonucleotide for human telomerase, BIOTIN P3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4.9%; Score 22; DB 1; Length 22; 100.0%; Pred. No. 1e+02; 1ve 0; Mismatches 0; Indels
                                                                                                                                                         Human clone 28-1 telomerase oligonucleotide oligo-P3.
                                                                                                                                                                                                                                                                                                                                           /*tag= a
/note= "5'-biotinylated cytidine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 22 BP; 3 A; 6 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Weinrich SL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure, Col 43-44; 34pp; English.
                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lichtsteiner SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ilarity 100.0%; P
Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP.
                                                 AAZ23628 standard; DNA; 22 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    97US-00833377.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        95US-00510736.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAS09473 standard; DNA; 22
                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1999-590379/50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
es 22; Conserv
                                                                                                                                                                                                                                                                                                         Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (GERO-) GERON
                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      04-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          04-AUG-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Atkinson EM,
Vasserot AP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24-OCT-2001
                                                                                                                         07-JAN-2000
                                                                                                                                                                                                                                                                                                                                                                                                US5968506-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                    19-0CT-1999
                                                                                                                                                                                                                  primer; ss
                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAS09473;
                                                                                     AAZ23628;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Loca
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAS09473/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 XEXEXEX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence represents an antisense oligonucleotide directed against the human telomerase RNA component gene sequences. The oligonucleotide can be used as an affinity agent in the methods of the invention, which are used to purify human telomerase. The methods of the use of several sequential steps, including the use of two matrixes that bind molecules bearing negative charges, a matrix that binds molecules bearing positive charges, an affinity purification step and a size separation. Telomerase is a particular target of anticancer therapies, and is useful in assays for characterizing (pre) cancerous cells. Telomerase can also be used to screen for specific modulators, for biochemical analysis of its activity, and in preparation of antibodies. Fragments of telomerase, or nucleic acid encoding them, are used in vaccines, and for treating over expression of telomerase, particularly in
ö
                                                                                                                                                                                                                                                                                                       Human; telomerase RNA component; anticancer therapy; purification; assay; vaccine; cancer; antisense oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pruzan RA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Purification of telomerase on affinity material - useful for, e.g. diagnosis and treatment of cancer.
                                                                                                                                                                                                                                                                           Antisense oligonucleotide P3 for human telomerase RNA component.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 22; DB 1; Length 22;
Pred. No. 1e+02;
0; Mismatches 0; Indels
 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 22 BP; 3 A; 6 C; 6 G; 7 T; 0 U; 0 Other;
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Atkinson EM, Lichtsteiner SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4.9%; Scor.
100.0%; Pred. No. 10.
0, Mismatches
 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                         /*tag= a
/note= "biotinylated"
                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                     163 GAGCAAACAAAAATGTCAGCT 184
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 24; 76pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            46 CTAACCCTAACTGAGAAGGGCG 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22 CTAACCCTAACTGAGAAGGGCG 1
                                                                    GAGCAAACAAAAATGTCAGCT
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                                                                                                                                                                  ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    97WO-US006012.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      97WO-US006012
                                                                                                                                                                 AAV63646 standard; DNA; 22
                                                                                                                                                                                                                                       (first entry)
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 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1998-594485/50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                         Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Weinrich SL,
                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    04-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        34-APR-1997;
                                                                                                                                                                                                                                         15-FEB-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9845450-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-0CT-1998.
   22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kealey JT;
                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local S:
Matches 22
                                                                                                                                                                                                     AAV63646;
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AAV63646//
IID AAV6.
IID AAV6.
XX AAC AAV6.
XX AAV6.
XX Human
XX H

Kealey JT;

Pruzan RA,

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Gaps

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WPI; 2003-403289/38.
                                                                   WO2003034985-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO2003034985-A2
                                               Homo sapiens
                                                                                                                                        22-OCT-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                          01-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-MAY-2003
                                                                                                                                                                                                            Ч,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ACC57543;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                           Rowley
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
 셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ò
                                                                                                                                                                                                                                                                                                                                                                                The sequence represents a biotinylated antisense oligomucleotide used in the purification of human telomerase. The invention relates to a purified human telomerase core enzyme protein comprising 2000-fold increased burners are protein comprising 2000-fold increased curvey compared with a crude extract of cells from adenovirus-transformed yearly compared with a crude extract of cells from adenovirus-transformed kidney cell line (193 cells) and when associated with telomerase RNA component has DNA polymerase activity and a molecular weight of 200-2000 kilo Daltons (RDA). The purified telomerase is useful for inducing a humoral or cell-mediated immune response in an animal. Purified telomerase or immunogenic fragments are useful as vaccines for treating diseases associated with over-expression of telomerase, such as cancer as affinity agents in isolating the proteins and for detecting the proteins and for the association of telomerase and/or telomerase associated proteins are also useful for screening compounds to identify agents that alter the association of telomerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                 Purified human telomerase, useful for inducing immune response in animals, comprises several thousand folds increased purity compared with
                                                                                                                                                                                                                                                         Pruzan RA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Telomerase; enzyme; RNA interference; short interfering RNA; siRNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
Human, Telomerase, vaccine, antibody, cancer; EF2H, nucleolin, antisense oligonucleotide, BIOTIN P3; ss.
                                                                                                                                                                                                                                                         Vasserot AP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4.9%; Score 22; DB 1; Length 22;
100.0%; Pred. No. 1e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 22 BP; 3 A; 6 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                         Lichtsteiner SP,
                                                                              /*tag= a
/mod_base= C
/note= "C is biotinylated"
                                                                                                                                                                                                                                                                                                                                         cytoplasmic crude cell preparations.
                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      46 CTAACCCTAACTGAGAAGGGCG 67
                                                                                                                                                                                                                                                                                                                                                                Disclosure; Col 18; 29pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CTAACCCTAACTGAGAAGGGCG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP
                                                                                                                                                                                               95US-00510736.
                                                                                                                                                                          99US-00420056
                                                                                                                                                                                                                                                          Atkinson EM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ACC57544 Standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 100.0
Warches 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Telomerase PCR primer.
                                                                                                                                                                                                                                                                                           WPI; 2001-450477/48.
                                                                                                                                                                                                                                  (GERO-) GERON CORP.
                                                        Key
modified_base
                                                                                                                                                                                                                                                     Weinrich SL,
                                    Homo sapiens
                                                                                                                            JS6261556-B1
                                                                                                                                                                          18-OCT-1999;
                                                                                                                                                                                                04-AUG-1995;
04-APR-1997;
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                                                                                                                                                   17-JUL-2001.
                                                                                                                                                                                                                                                                    Kealey JT;
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The present sequence is a PCR primer used for RT-PCR quantitation of telomerase RNA (see also ACC5751). The invention relates to the discovery that double-stranded interfering RNAs, such as short interfering RNAs (SIRNA), which target telomerase RNA or telomerase reverse transcriptase (TERN) mRNA, are capable of inhibiting telomerase activity. In cancer cells, inhibition of telomerase leads to telomere shortening, end-to-end chromosomal fusion, and apoptosis Telomerase inhibition can also be used for treatment of infertility, for contraception or sterilisation, for immunosuppression, for treatment of yeast, parasite and fungal infections, and in antiinflammatory therapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel nucleic acid encoding or comprising interfering RNAs which target telomerase RNA, useful for inhibiting telomerase activity for treating cancer, infertility and disorders of the immune system.
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cancer; tumour; cytostatic; contraceptive; immunosuppressive; antiinfertility; fungicide; antiparasitic; antiinflammatory; human; gene therapy; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Telomerase, enzyme; RNA interference; short interfering RNA; siRNA; cancer; tumour; cytostatic; contraceptive; immunosuppressive; antiinfertility; fungicide; antiparasitic; antiinflammatory; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
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100.0%; Pred. No. 1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 22 BP; 6 A; 7 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATGTCAGCTGCTGGCCCGTTCG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 2; Page 27; 52pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ACC57543 standard; DNA; 22 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-OCT-2002; 2002WO-US033146.
                                                                                                                                                                                                                                                                                                                                                                                                 16-OCT-2002; 2002WO-US033146.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2001US-0345326P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-FEB-2002; 2002US-0359196P.
22-MAY-2002; 2002US-0383195P.
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1es 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (UYRP ) UNIV ROCHESTER
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This invention relates to a purified human telomerase protein, which when associated with telomerase RNA component has DNA polymerase activity.

Also disclosed in the specification is a method for assessing a regulator (preferably a telomerase inhibitor or activator of telomerase involves measuring telomerase enzymatic activity of the composition in presence of a regulator. The telomerase protein of the invention may be used in the developing and testing assays for measuring telomerase activity which are useful in characterising cancer and pre-cancer cells, for identifying and testing aspainst telomerase. The mammalian telomerase protein of the invention is at least approximately 3000 fold more pure (in terms of telomerase activity per weight of protein) than a crude extract of the invention is at least approximately 3000 fold more pure (in terms of telomerase activity per weight of protein) than a crude extract of cell from adenovirus- transformed kidney cell. Purified telomerase (activity per weight of protein) than a crude extract of facilitates a thorough biochemical analysis of the enzyme's mechanism for developing mechanism-based regulators. The present sequence represents a human telomerase antisense oligonucleotide which has affinity to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Obtaining telomerase, by preparing enriched solution from cell expressing telomerase, combining the solution with oligonucleotide having specific affinity for the protein and collecting protein bound to oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                           telomerase and is used to purify the telomerase protein of the invention
Composition useful e.g. in diagnosis of cancer comprises complex of telomerase protein with telomerase RNA component.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mammalian telomerase protein; telomerase purification; telomere; anion exchange matrix; cation exchange matrix; selectivity matrix; gel filtration chromatography; gradient centrifugation; antisense oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human telomerase RNA component antisense oligonucleotide seq id
                                                                                                                                                                                                                                                                                                                                                                                                                         Score 22; DB 1; Length 22;
; Pred. No. 1e+02;
0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                             Sequence 22 BP; 3 A; 6 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lichtsteiner SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; SEQ ID NO 2; 24pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  46 CTAACCCTAACTGAGAGGGCG 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22 craacccraacreagaagggg 1
                                                       Disclosure; Col 9; 24pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            95US-00510736.
97US-00833377.
99US-00420056.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ВР.
                                                                                                                                                                                                                                                                                                                                                                                                                                               100.08;
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                                                                                                                                                                                                                                                                                                                                                                                                                                4.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADC35649 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                               Ouery Match
Best Local Similarity luve.
Fra 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Atkinson
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Weinrich SL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US6545133-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-APR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              08-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADC35649;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADC35649,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                          interfering RNAS (SIRNA), which target telomerase RNA or telomerase reverse transcriptase (TERT) mRNA, are capable of inhibiting telomerase activity. In cancer cells, inhibition of telomerase leads to telomerase hortening, end-to-end chromosomal fusion, and apoptosis. Telomerase inhibition can also be used for treatment of infertility, for contraception or sterilisation, for immunosuppression, for treatment of yeast, parasite and fungal infections, and in antiinflammatory therapies
                                                                                                                                                                            Novel nucleic acid encoding or comprising interfering RNAs which target telomerase RNA, useful for inhibiting telomerase activity for treating cancer, infertility and disorders of the immune system.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pruzan RA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                             The present sequence is a PCR primer used for RT-PCR quantitation of telomerase RNA (see also ACC57551). The invention relates to the discovery that double-stranded interfering RNAs, such as short
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.9%; Score 22; DB 1; Length 22;
.00.0%; Pred. No. 1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human telomerase antisense oligonucleotide primer P3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 22 BP; 2 A; 3 C; 11 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Telomerase; antisense; primer; P3; ss; cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Atkinson EM, Lichtsteiner SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /*tag= a
/note= "Biotinylated"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CTGGGAGGGGTGGTGGCCATTT 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CTGGGAGGGGTGGTGGCCATTT 22
                                                                                                                                                                                                                                                 Example 2; Page 27; 52pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ВР
        22-OCT-2001; 2001US-0345326P.
20-FEB-2002; 2002US-0359196P.
22-MAY-2002; 2002US-0383195P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.08;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     97US-00833377
99US-00420056
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABX10983 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22; Conservative
                                                                              (UYRP ) UNIV ROCHESTER
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                                                                                                                                            WPI; 2003-403289/38
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Weinrich SL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        04-AUG-1995;
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18-OCT-1999;
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                                                                                                               Rowley PT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19
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Gaps

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RA;

Pruzan

Vasserot AP,

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The invention describes a method of obtaining mammalian telomerase protein (I). The method involves preparing enriched solution (ES) from a cell expressing telomerase where the component of (I) in ES is separated from other proteins expressed by cell by combining ES with coligonuclectide (O) having specific affinity for (I), and collecting protein bound to (O). The oligonuclectide comprises a retrievable label such as biotin and contains a sequence that is specifically recognized by telomerase protein. The oligonuclectide contains or does not contain the sequence (TTAGGG). The method further comprises combining a fraction containing telomerase protein with a maion exchange matrix, and collecting protein with a cation exchange matrix. (such as a heparin matrix), and collecting protein that binds the matrix. The method containing telomerase protein that binds the matrix. The method on several different ion exchange matrices and combining a fraction containing telomerase protein with an intermediate selectivity matrix, collecting protein that binds the matrix, where the intermediate selectivity matrix, and separating a fraction containing telomerase protein with an intermediate selectivity matrix, collecting protein that binds the matrix, where the intermediate selectivity matrix, and separating a fraction containing the telomerase is enriched from an extract of cells stably expressing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                                                              telomerase. This sequence represents an antisense oligonucleotide to the
                                                                                                                                                                                                                                                                                                                                                                                               component of human telomerase that can be used in the purification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pruzan RA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Telomerase activity; therapy; cancer; cytostatic; antisense; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Lichtsteiner SP, Vasserot AP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4.9%; Score 22; DB 1; Length 22;
100.0%; Pred. No. 1e+02;
ve 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human telomerase RNA antisense oligonucleotide, P3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Seguence 22 BP; 3 A; 6 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "Biotin labelled"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               46 CTAACCCTAACTGAGAAGGCG 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CTAACCCTAACTGAGAAGGGCG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ig= a
|_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; Fre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADG62871 standard; DNA; 22 BP.
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20-NOV-2000; 2000US-00717828.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (WEIN/) WEINRICH S L.
(ATKI/) ATKINSON E M.
(LICH/) LICHTSTEINER S P.
(VASS/) VASSEROT A P.
(PRUZ/) PRUZAN R A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Weinrich SL, Atkinson EM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                  method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
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04-APR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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ADG62871/c
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                                                                                                              The invention relates to a method for identifying regulators of telomerase activity that may be useful for treating cancers. The method may be used to identify regulators e.g. antibodies, of telomerase activity which may be useful as cancer treatments. It has been found that found that the cells of many human cancers have telomerase activity. This sensement. If telomerase activity in cancer cells can be inhibited, the cancer cells are expected to reach sensemence and cease dividing. The present sequence is human telomerase antisense oligonulceotide used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Telomerase; enzyme; RNA interference; short interfering RNA; siRNA; telomerase; cancer; tumour; cytostatic; contraceptive; immunosuppressive; antiinfertility; fungicide; antiparasitic; antiinflammatory; human; gene therapy; PCR; primer; sa.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleic acids which are complementary to each other and to target nucleic acid e.g., telomerase RNA or mRNA encoding telomerase reverse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel double-stranded short interfering RNA having sense and antisense
                                                                                                                                                                                                                                                                                                                                                                        Gapa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence is a PCR primer used for RT-PCR quantitation of telomerase RNA (see also ACC58040). The invention relates to the discovery that double-stranded interfering RNAs, such as short interfering RNAs (SIRNA), which target telomerase RNA or telomerase
                                                                                                                                                                                                                                                                                                                                                                        ö
                                                 Identifying telomerase regulators useful for treating cancer.
                                                                                                                                                                                                                                                                                                                                    4.9%; Score 22; DB 1; Length 22;
100.0%; Pred. No. 1e+02;
                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                  Sequence 22 BP; 3 A; 6 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                     illustrate the method of the invention.
                                                                                Disclosure; SEQ ID NO 2; 22pp; English
                                                                                                                                                                                                                                                                                                                                                                                                        46 CTAACCCTAACTGAGAAGGGCG 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 2; Page 24; 37pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                             CTAACCCTAACTGAGAAGGGCG 1
                                                                                                                                                                                                                                                                                                                                                     100.0%;
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20-FEB-2002; 2002US-0359196P.
22-MAY-2002; 2002US-0383195P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ACC58032 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            relomerase PCR primer.
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                                                                                                                                                                                                                                                                                                                                                     Local Similarity
nes 22; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO2003035667-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACC58032;
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22;

Matches

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               activity. In cancer cells, inhibition of telomerase leads to telomere shortening, end-to-end chromosomal fusion, and apoptosis. Telomerase inhibition can also be used for treatment of infertility, for contraception or sterilisation, for immunosuppression, for treatment of yeast, parasite and fungal infections, and in antiinflammatory therapies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  telomerase; cancer; tumour; cytostatic; contraceptive; immunosuppressive; antiinfertility; fungicide; antiparasitic; antiinflammatory; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel double-stranded short interfering RNA having sense and antisense unducleic acids which are complementary to each other and to target nucleic acid e.g., telomerase RNA or mRNA encoding telomerase reverse transcriptase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence is a PCR primer used for RT-PCR quantitation of telomerase RNA (see also ACC58040). The invention relates to the discovery that double-grranded interfering RNAs, such as short interfering RNAs, such as short interfering RNAs (sIRNA), which target telomerase RNA or telomerase reverse transcriptase (TERT) and EAD of telomerase reverse transcriptase (TERT) inhibition of telomerase leads to telomerase shortening, end-to-end chromosomal fusion, and apoptosis Telomerase inhibition can also be used for treatment of infertility, for contraception or sterilisation, for immunosuppression, for treatment of yeast, parasite and fungal infections, and in antiinflammatory therapies
are capable of inhibiting telomerase
                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Telomerase, enzyme, RNA interference, short interfering RNA, siRNA,
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                                                                                                                                                                    DB 1; Length 22;
1e+02;
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                                                                                                                            Sequence 22 BP; 6 A; 7 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                  4.9%; Score 22; DB ilarity 100.0%; Pred. No. 1e+ Conservative 0; Mismatches
                                                                                                                                                                                                                                       176 ATGTCAGCTGCTGGCCGTTCG 197
reverse transcriptase (TERT) mRNA,
                                                                                                                                                                                                                                                                           22 ArGrcAGCrGCrGGCCCGrrcG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gene therapy; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2001US-0345326P
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2002US-0383195P.
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                                                                                                                                                                                                                                                                                                                                                                   ACC58031 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Telomerase PCR primer.
                                                                                                                                                                Query Match
Best Local Similarity
Matches 22; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-403336/38.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22-OCT-2001;
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                                                                                                                                                                                                                                                                                                                                 RESULT 129
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    mouse, rat and chinese master are all claimed. Antisense oligonucleotides can be used to block the activity of the telomerase; probes and primers can be used to block the activity of the telomerase; probes and primers can be used in detection; vectors and host cells transformed with the isolated telomerase genes can be used for production of telomerases; RNA and DNA ribozymes and triplex forming of ignoricectides directed against the telomerase genes can be used therapeutically as can plasmids. A mouse which lacks the telomerase gene (also claimed) can be used for study of telomere regulation in vivo, and AATI1059, AATI1069, which are complementary to human telomerase RNA component sequences can be used to identify and amplify homologous sequences from other non-human mammals. The amplified fragments can then be used as probes to identify telomerase genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   components of (non)human mammalian telomerase(s) - useful in studying
                                                                                                                                                                                                                                                Telomerase; mammal; antisense; triplex forming oligonucleotide; plasmid;
      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Marhuenda MA;
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0
    Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Greider C,
                                                                                                                                                                                                                    Primer used for amplifying telomerase RNA fragments.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21 BP; 5 A; 5 C; 11 G; 0 T; 0 U; 0 Other;
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0
 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Funk W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Pred. ....
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RNA components of (non)human mammalia cell senescence and immortalisation.
                              40
                                                        22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 15; Page 59; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Feng J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  184 TGCTGGCCCGTTCGCCCTCC 204
                                                        CTGGGAGGGGTGGTGGCCATTT
 ;
                             19 CTGGGAGGGGTGGTGGCCATTT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (COLD-) COLD SPRING HARBOR LAB
                                                                                                                                   BP.
                                                                                                                                                                                                                                                                                                                                                                                                          94US-00272102.
94US-00330123.
95US-00387524.
95US-00485778.
                                                                                                                                                                                                                                                                                                                                                                                95WO-US008620
                                                                                                                                                                                                                                                               probe; primer; ribozyme; ss.
                                                                                                                             AAT11058 standard; DNA; 21
                                                                                                                                                                                       (first entry)
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Avillon AA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1996-097428/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GERO-) GERON CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
Les 21; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Villeponteau B;
                                                                                                                                                                                                                                                                                                                      WO9601614-A2
                                                                                                                                                                                                                                                                                                                                                                                07-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                          07-JUL-1994;
27-OCT-1994;
                                                                                                                                                                                       02-JUL-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                   25-JAN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Andrews WH,
                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                           AAT11058;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                               Best Loca
Matches
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RESULT 131

4.9%; Score 22; DB 1; Length 22; 100.0%; Pred. No. 1e+02;

Query Match Best Local Similarity

96WO-US009517. 95US-00478352.

Autexier C;

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DNA encoding essential RNA components of human telomerase - also truncated or recombinant telomerase, useful for diagnosis and treatment of cancer and infection by eukaryotic parasites.
                                                                                                                            (COLD-) COLD SPRING HARBOR LAB
                                                                                                                                                                           WPI; 1997-099928/09
                                                                                                      07-JUN-1995;
                                  WO9640868-A1
                                                         19-DEC-1996.
                                                                                                                                                     Greider C,
           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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셤
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence is a PCR primer for the RNA component of a non-human mammal telomerase (RCT), cDNA. The RCT can be used in the recombinant prodn. of an active telomerase mol., capable of adding sequences to chromosomal DNA telomers, and in the synthesis of mutant sequences for the detection of mutant mammalian telomerase RNA component polynucleotides. The RCT may also be used in the identification of telomerase modulating agents, and in the detection of telomerase related, or neoplastic conditions in a patient. Polynucleotides of at least 25 consecutive nucleotides identical, or complementary to the RCT sequence linked to heterologous transcriptional regulatory sequences, can be used for the gene therapy of human diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cancer, eukaryotic parasite, hTR; vertebrate telomerase, yeast; protozoa, tumour; antibody; polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                       RNA component of mammalian telomerase, esp. human - useful in identifying e.g. candidate telomerase-modulating agents.
                                                                                                               detection; mammalian; identification; modulating agent; neoplastic condition; transcriptional regulatory sequence; gene therapy; disease; polymerase chain reaction; PCR primer; non-human; ss.
                                                                                                        component; telomerase; recombinant production; synthesis; mutant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                RNA component of non-human mammal telomerase cDNA PCR primer R7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4.7%; Score 21; DB 1; Length 21; 100.0%; Pred. No. 1.2e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 5 A; 5 C; 11 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                             Funk W, Andrews WH;
                                                                                                                                                                                                                                                                                                                                                                                                                             Example 9; Page 81; 114pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  184 TGCTGGCCCGTTCGCCCCTCC 204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human telomerase PCR 3'-primer R7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TGCTGGCCCGTTCGCCCCTCC 1
           AAT10301 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                      94US-00330123.
95US-00472802.
95US-00482115.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAT58813 standard; DNA; 21 BP
                                                                                                                                                                                                                                    95WO-US008530
                                                                                                                                                                                                                                                             94US-00272102
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                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                             Villeponteau B, Feng J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21; Conservative
                                                                                                                                                                                                                                                                                                                                                                    WPI; 1996-097581/10.
                                                                                                                                                                                                                                                                                                                     (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                      WO9601835-A1.
                                                                                                                                                                                                                                   06-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-NOV-1997
                                                         10-SEP-1996
                                                                                                                                                                                                                                                                        27-OCT-1994;
07-JUN-1995;
                                                                                                                                                                                                             25-JAN-1996
                                                                                                                                                                                                                                                                                              07-JUN-1995
                                                                                                                                                               Synthetic.
                                  AAT10301;
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AAT10301/c
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                                                                                                        RNA
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The present sequence represents PCR 3'-primer R7 used for amplifying the human telomerase (hTR). The RNA and DNA can be used in hybridisation cassays to detect or quantify telomerase activity in cells, tissue or fluid samples, e.g. for diagnosis of eukaryotic parasites (yeast and protozoa) or tumours. It is also useful as primers for amplification or assays. The truncated or recombinant vertebrate telomerase is used the charapeutically to increase telomerase activity (also as reagents in the screening assay) while the RNA or other inhibitors such as antisense of therapeutically to reduce such activity. Typical applications are initiation/restoration of activity to cause senescence or to prevent immortalisation of cells in tumours or parasites. The DNA is also used to produce recombinant telomerase, which can then be used conventionally to raise antibodies for diagnostic detection of telomerase. Detecting telomerase antibodies for diagnostic detection of telomerase. Detecting telomerase antibodies are diagnosis of tumour or infection, before clinical signs annifest. Telomerase inhibitors directed against e.g. Trypanosoma should cause fewer side effects than drugs currently used to treat such infections. The DNA encodes those parts of hTR RNA assential for activity infections. The DNA encodes those parts of hTR RNA assential for activity infections.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             neuroprotective, anti-HIV; ophthalmological; antiulcer; antirhoumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; shNA; shNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 21 BP; 5 A; 5 C; 11 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pred. No. 1.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Pred. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           184 TGCTGGCCCGTTCGCCCTCC 204
Example 5; Page 32; 48pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADF93862 standard; RNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human TERT siRNA, SEQ ID 589.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26-FEB-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADF93862;
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Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short interfering RNA; shraw; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.

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New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                             Seguence 21 BP; 2 A; 7 C; 8 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                      Example 3; SEQ ID NO 589; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           397 GCGCCCGATTCCCTGAGCTG 417
                                                                          06-UIN-2002; 2002US-0386782P.
17-JUL-2002; 2002US-0396600P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
                                                                                                          09-SEP-2002; 2002US-0409293P.
                                                                                                                                                                                                                                                                                                                                                                                                                            4.7%;
81.0%;
                                             11-FEB-2003; 2003WO-US004088
                                                                                                                                  (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                 Mcswiggen J, Beigelman L;
                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 81.0%
                                                                                                                                                                WPI; 2003-689777/65.
               WO2003070742-A1
Homo sapiens.
                                                            20-FEB-2002;
11-MAR-2002;
                              28-AUG-2003.
셤
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New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.

2002US-0363124P. 2002US-0386782P. 2002US-0396600P.

06-JUN-2002; 17-JUL-2002; 29-AUG-2002;

11-MAR-2002; 20-FEB-2002;

11-FEB-2003; 2003WO-US004088

WO2003070742-A1 Homo sapiens.

Synthetic

28-AUG-2003

2002US-0406784P. 2002US-0408378P. 2002US-0409293P.

05-SEP-2002; 15-JAN-2003;

2003US-0440129P

(RIBO-) RIBOZYME PHARM INC. Mcswiggen J, Beigelman L;

WPI; 2003-689777/65.

ö downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration. screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents a siRNA targeted to the human TERT mRNA transcript. skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, invention relates to short interfering nucleic acids (siNA) which Gaps ·. Score 21; DB 1; Length 21; Pred. No. 1.2e+02; 4; Mismatches 0; Indels GCGCCCGAUUCCCUGAGCUG 21 Human TERT siRNA, SEQ ID 594. ADF93867 standard; RNA; 21 (first entry) 26-FEB-2004 ADF93867; RESULT 134 ADF93867/C 1D ADF93 XX AC ADF93 XX DT 26-FF XX XX XX XX XX XX

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The invention relates to short interfering nucleic acids (sinA) which downregulate expression of the one or more telomerase genes by RNA interference. The sinAs may or may not comprise shoul electides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an misense oligonucleotide. Specifically, the sinAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The sinAs can be unmodified or chemically modified, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restencisis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lubus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             engineering, pharmacogenomics, studying gene function, and gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 3 A; 8 C; 8 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 3; SEQ ID NO 594; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             395 GCGCGCGCGATTCCCTGAGC 415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21 GCGCGCGCGATTCCCTGAGC 1
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Best Local Similarity
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à
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Cytostatic, vasotropic, protozoacide, immunosuppressive, dermatological; neuroprotective, anti-HIV, ophthalmological, antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase, human, terc; RNA interference, shour interfering nucleic acid; siNA, short interfering NNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 3; SEQ ID NO 587; 145pp; English.
                     ADF93860 standard; RNA; 21 BP
                                                                                                                                                                                                                                                                                                                                 2002US-0396600P.
2002US-0406784P.
2002US-0408378P.
                                                                                  Human TERT siRNA, SEQ ID 587.
                                                                                                                                                                                                                                                                                                   2002US-0358580P.
2002US-0363124P.
2002US-0386782P.
                                                                                                                                                                                                                                                                                                                                                                 2002US-0409293P
2003US-0440129P
                                                                                                                                                                                                                                                                             11-FEB-2003; 2003WO-US004088
                                                                                                                                                                                                                                                                                                                                                                                                 (RIBO-) RIBOZYME PHARM INC.
                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                     Mcswiggen J, Beigelman L;
                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-689777/65.
                                                                                                                                                                                                                                   WO2003070742-A1.
                                                                                                                                                                                                                                                                                                                                  17-JUL-2002; 29-AUG-2002; 2
                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                 09-SEP-2002;
                                                                                                                                                                                                                                                                                                    20-FEB-2002;
                                                                                                                                                                                                                                                                                                                        06-JUN-2002;
                                                             26-FEB-2004
                                                                                                                                                                                                                                                         28-AUG-2003
                                                                                                                                                                                                     Synthetic
                                         ADF93860;
RESULT 135
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclecides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonuclectide and an interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modilate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AlbS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug cancering, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents a siRNA targeted to the human TERT mRNA transcript.

Sequence 21 BP; 2 A; 10 C; 3 G; 0 T; 6 U; 0 Other;

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonuclectide and an antisense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairbin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised, and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be
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                                                                           Gaps
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Score 21; DB 1; Length 21;
Pred. No. 1.2e+02;
6; Mismatches 0; Indels
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                                                                                                                                              138 CTGCCGCCTTCCACCGTTCAT 158
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2002US-0406784P.
2002US-0408378P.
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2002US-0386782P.
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4.7%;
ilarity 71.4%;
Conservative
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   Query Match
Best Local Similarity
Matches 15; Conserv
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17-JUL-2002;
29-AUG-2002;
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used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, es; multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleoide polymorphisms). The present sequence represents a siRNA targeted to the human TERT mRNA transcript.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
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Pred. No. 1.2e+02;
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Best Local Similarity 100.0%; Pred. No. 1.26
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11-WAR-2002; 2002US-0363124P.

06-UUN-2002; 2002US-0386782P.

17-UUL-2002; 2002US-0396600P.

29-AUG-2002; 2002US-0406784P.

05-SEP-2002; 2002US-0409378P.

15-JAN-2003; 2003US-0401299P.
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ADP939866/

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an antisense oligonucleotide. Specifically, the siNAs include short
interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short
can contain Geoxyribonucleotides, and can be chemically wondified,
can contain deoxyribonucleotides, and can be chemically withesised,
can contain deoxyribonucleotides, and can be chemically synthesised,
cand/or complexes to kits for the in vitro or in vivo delivery of siNA, conjugates
and/or complexes of siNA, and vectors that express siNA. The siNAs are
cused to modulate expression of the telomerase genes in cells, tissue
cused to modulate expression of the telomerase genes in cells, tissue
cused to modulate expression of the telomerase genes in cells, tissue
cused to modulate refection of avariety of conditions. They may be
transplants for the treatment of a variety of conditions. They may be
cused for treating cancer, restenosis, infectious diseases (specifically
protozoal), transplant rejection, or autoimmune or age-related diseases,
c.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration,
skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug
correction, displandicion and validation,
genetic engineering, pharmacogenomics, studying gene function, and gene
cmapping (e.g., of single nucleotide polymorphisms). The present sequence
ceptivesents a siRNA targeted to the human TERT mRNA transcript.
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2002US-0363124P.
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2002US-0396600P.
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2002US-0408378P.
2002US-0409293P.
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06-JUN-2002; 2
17-JUL-2002; 2
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense cregions, or alternatively are assembled from a sense oligonuclectide and antisense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short contain deoxyribonuclectides, and can be unmodified or chemically synthesised, can contain deoxyribonuclectides, and can be chemically synthesised, can contain deoxyribonuclectides, and can be chemically synthesised, can contain deoxyribonuclectides, and can be chemically synthesised. Cc expressed from a vector or renzymatically synthesised. The invention also crelates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express sincells, tissue cexplants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene companies a siRNA targeted to the human TERT mRNA transcript.
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New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                           Example 3; SEQ ID NO 586; 145pp; English.
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2002US-0386782P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-FEB-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 81.0
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO2003070742-A1.
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11-MAR-2002; 2
06-JUN-2002; 2
17-JUL-2002; 2
29-AUG-2002; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 139
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The invention relates to short interfering nucleic acids (sINA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue cared to modulate expression of the telomerase genes in cells, tissue used for transplants for the treatment of a variety of conditions. They may be used for transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents a siRNA targeted to the human TERT mRNA transcript.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                 New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Mismatches
                                                                                                                                                                                                                                                                                                   Example 3; SEQ ID NO 588; 145pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 ACCCACUGCCACCGCGAAGAG 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADF93865 standard; RNA; 21 BP
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human TERT siRNA, SEQ ID 592.
                                                                                            (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                         Mcswiggen J, Beigelman L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-FEB-2004 (first entry)
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Atches 20; Conservative
                                                                                                                                                                                      WPI; 2003-689777/65.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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ADF93865/c
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epilepsy, dementia; huntington's;
Alzheimer's; Parkinson's; epilepsy; dementia; huntin
amyotrophic lateral sclerosis; gene therapy; ss; hTR
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Jamison S, Usman N, Thompson J;
                                                                                                                                                                                                                                                                                                       (SIRN-) SIRNA THERAPEUTICS INC
                                                                                                                                                                                                                     2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
                                                                                                                                                                                         2002US-0363124P.
2002US-0386782P.
                                                                                                                                                                          2002US-0358580P.
                                                                                                                                                                                                                                                                       15-JAN-2003; 2003US-0440129P.
                                                                                                                                            20-FEB-2003; 2003WO-US005028
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                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-731676/69.
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                                                                              WO2003074654-A2
                                                                                                                                                                           20-FEB-2002;
                                                                                                                                                                                                                          29-AUG-2002;
05-SEP-2002;
                                                                                                                                                                                                                                                        09-SEP-2002;
                                                                                                                                                                                                           06-JUN-2002;
                                                                                                              12-SEP-2003
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                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense or regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short can contain deoxyribonucleotides, and can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue cyplants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically processed), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single mucleotide polymorphisms). The present sequence mapping (e.g., of single mucleotide polymorphisms). The present sequence creaming a siRNA targeted to the human TERT mRNA transcript.
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                                                                                                                                                                                                                                                                                                                      New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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Pred. No. 1.2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              double-stranded short interfering nucleic acid; siNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 5 A; 4 C; 11 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                      Example 3; SEQ ID NO 592; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GCCTGCCGCCTTCCACCGTTC 156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hTR-targeted sina RNA - SEQ ID 597
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Local Similarity 100.0%; Pr
                                                                              20-FEB-2002; 2002US-0358580P.
11-MAX-2002; 2002US-0363124P.
06-UIN-2002; 2002US-0386782P.
17-UUL-2002; 2002US-0396600P.
29-AUG-2002; 2002US-0406784P.
09-SEP-2002; 2002US-0409378P.
                                                                                                                                                                                            LS-JAN-2003; 2003US-0440129P.
                                                  11-FEB-2003; 2003WO-US004088
                                                                                                                                                                                                                            (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADG30031 standard; RNA; 21
                                                                                                                                                                                                                                                            Mcswiggen J, Beigelman L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                          WPI; 2003-689777/65.
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                  28-AUG-2003
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Chowrira B, Pavco P, Fosnaugh K;

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The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of the invention demonstrates antiarteriosclerotic, neuroprotective, noctropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary diseases such as restenosis, atherosclerosis, Alzheimer's disease, Parkinson's disease, epilepsy, dementia, huntington's disease, epilepsy, dementia, huntington's disease or amyotrophic lateral sclerosis. Furthermore, the sinA may be utilised for gene, therapy applications. The current sequence is that of the siNA RNA
New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21 BP; 2 A; 7 C; 8 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4; Mismatches
                                                                                                                                                                                                                                                                     Example 24; SEQ ID NO 597; 593pp; English.
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The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of the invention demonstrates antiarteriosalerotic, neuroprotective, nootropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           diseases such as restenosis, atherosclerosis, Aîzheimer's disease, Parkinson's disease, epilepsy, dementia, huntington's disease or amyotrophic lateral sclerosis. Purthermore, the siNA may be utilised for gene therapy applications. The current sequence is that of the siNA RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antiarteriosclerotic; neuroprotective; nootropic; antiparkinsonian; anticonvulsant; pulmonary disease; restenosis; atherosclerosis;
  Alzheimer's; Parkinson's; epilepsy; dementia; huntington's; amyotrophic lateral sclerosis; gene therapy; se; hTR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4.7%; Score 21; DB 1; Length 21; 95.2%; Pred. No. 1.2e+02; Artive 1; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 21 BP; 6 A; 9 C; 5 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 24; SEQ ID NO 596; 593pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chowrira B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hTR-targeted sina RNA - SEQ ID 602.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       285 ACCCACTGCCACCGCGAAGAG 305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 ACCCACUGCCACCGCGAAGAG 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Thompson J;
                                                                                                                                                                                                                                                                                                      2002US-0363124P.
2002US-0386782P.
2002US-0406784P.
2002US-0408378P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (SIRN-) SIRNA THERAPEUTICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP.
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15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                                 20-FEB-2003; 2003WO-US005028
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Beigelman L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADG30036/c
ID ADG30036 standard; RNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 J, Bers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-731676/69.
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Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     of the invention
                                                                                                                            WO2003074654-A2
                                                                             Unidentified
                                                                                                                                                                                                                                                                                    20-FEB-2002;
                                                                                                                                                                                                                                                                                                         11-MAR-2002;
                                                                                                                                                                                                                                                                                                                                      06-JUN-2002;
                                                                                                                                                                                                                                                                                                                                                         29-AUG-2002;
05-SEP-2002;
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                                                                                                                                                                               12-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Jamison S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mcswiggen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  activity.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a double-stranded short interfering nucleic acid (sinA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded sinA comprises about 21 nucleotides. The sinA of the invention demonstrates antiarteriosclerotic, neuroprotective, nootropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian thar responds to endilation of gene expression or activity in a cell, tissue or organism. The disease or condition any include pulmonary diseases such as restenosis, atherosclerosis, Alzheimer's disease, parkinson's disease, palleppy, dementia, hutington's disease or amyotrophic lateral sclerosis. Furthermore, the sinA may be utilised for gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or
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  dementia; huntington's;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
4.7%; Score 21; DB 1; Length 21;
Best Local Similarity 81.0%; Pred. No. 1.2e+02;
Matches 17; Conservative 4; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             double-stranded short interfering nucleic acid; siNA;
Alzheimer's; Parkinson's; epilepsy; dementia; huntin
amyotrophic lateral sclerosis; gene therapy; ss; hTR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 21 BP; 2 A; 3 C; 12 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 24; SEQ ID NO 594; 593pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 UUGCGGAGGGUGGGCCUGGGA 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Beigelman L, Chowr
man N, Thompson J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (SIRN-) SIRNA THERAPEUTICS INC
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                                                                                                                                                                                                                                                                                                            2002US-0363124P.
2002US-0386782P.
                                                                                                                                                                                                                                                                                                                                                      29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408376P.
09-SEP-2002; 2002US-0409293P.
                                                                                                                                                                                                                                                                                                                                                                                                                                        15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                               20-FEB-2003; 2003WO-US005028
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                                                                                                                            WO2003074654-A2
                                                                          Unidentified.
                                                                                                                                                                                                                                                                                                                                   06-JUN-2002;
                                                                                                                                                                                                                                                                                                            11-MAR-2002;
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Foenaugh K;

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Alzheimer's; Parkinson's; epilepsy; dementia; huntington's; amyotrophic lateral sclerosis; gene therapy; ss; hTR.
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2002US-0386782P.
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2002US-0408378P.
2002US-0409293P.
                                                                                                                2003WO-US005028
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                                                                                                                                                                                                                                                                                                   WPI; 2003-731676/69.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                              WO2003074654-A2
                                                                                                                                                                           29-AUG-2002; 2
05-SEP-2002; 2
09-SEP-2002; 2
15-JAN-2003; 2
                                                                                                                20-FEB-2003;
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                                                                                                                                                                                                                                                                                                                                                                  activity.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                the invention demonstrates and artiatreniosclerotic, neuroprotective, nootropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary diseases such as restenosis, atherosclerosis, Albrahimer's disease, Parkinson's disease, epilepsy, dementia, huntington's disease or amyotrophic lateral sclerosis. Furthermore, the siNA may be utilised for gene therapy applications. The current sequence is that of the siNA RNA
                                                                                                                                                                                                                                                                                                                            New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or activity.
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0
                                                                                                                                                                                                                                                                  Pavco P, Fosnaugh K;
  Alzheimer's, Parkinson's; epilepsy; dementia; huntington's; amyotrophic lateral sclerosis; gene therapy; ss; hTR.
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Pred. No. 1.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 3 A; 8 C; 8 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                           Example 24; SEQ ID NO 602; 593pp; English.
                                                                                                                                                                                                                                                                 Beigelman L, Chowrira B,
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                                                                                                                                                                                                                                                                               Thompson J;
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                                                                                                                                                   2002US-0363124P.
2002US-0386782P.
2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
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                                                                                                                                         2002US-0358580P.
                                                                                                                 20-FEB-2003; 2003WO-US005028
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                                                                                                                                                                                                                                                                              Usman N,
                                                                                                                                                                                                                                                                                                    WPI; 2003-731676/69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the invention
                                                                 WO2003074654-A2
                                                                                                                                                                                         05-SEP-2002;
09-SEP-2002;
                                                                                                                                                                                                                   L5-JAN-2003;
                                         Unidentified
                                                                                                                                                      11-MAR-2002;
                                                                                                                                                                06-JUN-2002;
                                                                                                                                                                               29-AUG-2002;
                                                                                                                                                                                                                                                                    Mcswiggen J,
                                                                                                                                          20-FEB-2002;
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                                                                                          12-SEP-2003
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                                                                                                                                                                                                                                                                               Jamison S,
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(sinA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded sinA comprises about 21 nucleotides. The sinAn of the invention demonstrates antiarteriosclerotic, neuroprotective, nootropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary diseases such as restenosis, atherosolerosis, Albrimer's disease, Parkinson's disease, epilepsy, dementia, huntington's disease or amyotrophic lateral sclerosis. Furthermore, the siNA may be utilised for gene therapy applications. The current sequence is that of the siNA RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a double-stranded short interfering nucleic acid
                                                                                                                                                                                                                                                                                                             New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ó
                                                                                     Chowrira B, Pavco P, Fosnaugh K;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 24; SEQ ID NO 601; 593pp; English.
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                                                                                     Mcswiggen J, Beigelman L, Chowri
Jamison S, Usman N, Thompson J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21 GCACCCACTGCCACCGCGAAG 1
(SIRN-) SIRNA THERAPEUTICS INC
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Alzheimer's; Parkinson's; epilepsy; dementia; huntington's; amyotrophic lateral sclerosis; gene therapy; ss; hTR.
                                                              WO2003074654-A2
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06-JUN-2002;
29-AUG-2002;
05-SEP-2002;
                                      Unidentified
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Local Si...
21;
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                                                                                                                                                                                                                                                                                     Jamison S,
                                                                                                                                                                                                                                                                         Mcswiggen
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                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene compitising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of the invention demonstrates antlarteriosclerotic, neuroprotective, notropic, antiparkineonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary diseases such as restenosis, atherosclerosis, Alzheimer's disease, Parkinson's disease, epilepsy, dementia, huntington's disease or amyotrophic lateral sclerosis. Furthermore, the siNA may be utilised for
                                                                                                                                                                                                                                                                                                                                   New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene therapy applications. The current sequence is that of the siNA RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                       Chowrira B, Pavco P, Fosnaugh
 Alzheimer's; Parkinson's; epilepsy; dementia; huntington's; amyotrophic lateral sclerosis; gene therapy; 88; hTR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21 BP; 2 A; 10 C; 3 G; 0 T; 6 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 21; DB 1; I
Pred. No. 1.2e+02;
6; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                      Example 24; SEQ ID NO 595; 593pp; English.
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| CUGCCGCCUUCCACCGUUCAU 21
                                                                                                                                                                                                                                                                                     Thompson J;
                                                                                                                                        2002US-0358580P.
2002US-0363124P.
2002US-0386782P.
2002US-0406784P.
2002US-0408378P.
                                                                                                                                                                                                                                               (SIRN-) SIRNA THERAPEUTICS INC
                                                                                                                                                                                                                      15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                       2002US-0409293P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4.7%;
ilarity 71.4%;
Conservative
                                                                                                                 20-FEB-2003; 2003WO-US005028
                                                                                                                                                                                                                                                                         Beigelman L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADG30033 standard; RNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                            Mcswiggen J, Belyerman S. Usman N,
                                                                                                                                                                                                                                                                                                           WPI; 2003-731676/69.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
ses 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the invention
                                                                WO2003074654-A2
                                        Unidentified
                                                                                                                                                                                29-AUG-2002;
05-SEP-2002;
                                                                                                                                                                                                        09-SEP-2002;
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                                                                                        12-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                            activity.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 147
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The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of the invention demonstrates antiarteriosclerotic, neuroprotective, nootropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     diseases such as restenosis, atheroscierosis, Alzheimer's disease, Parkinson's disease, epilepsy, dementia, huntington's disease or amyotrophic lateral sclerosis. Furthermore, the siNA may be utilised for gene therapy applications. The current sequence is that of the siNA RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          double-stranded short interfering nucleic acid; siNA; antiarteriosclerotic; neuroprotective; nootropic; antiparkinsonian; anticonvulsant; pulmonary disease; restenosis; atherosclerosis;
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                                                                                                                                                                                                                                                                                                                                        Fosnaugh
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Pred. No. 1.2e+02;
0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                        B, Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21 BP; 4 A; 13 C; 3 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 24; SEQ ID NO 599; 593pp; English.
                                                                                                                                                                                                                                                                                                                                        Chowrira
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         hTR-targeted sina RNA - SEQ ID 600.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GGTTGCGGAGGGTGGGCCTGG 22
                                                                                                                                                                                                                                                                                                                                                                     Thompson J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Pre-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GGTTGCGGAGGGTGGGCCTGG 1
                                                                                                                                                                                                                                                                                   (SIRN-) SIRNA THERAPEUTICS INC
                                                                              2002US-0363124P.
2002US-0386782P.
2002US-0406784P.
2002US-0408378P.
                                                                                                                                                                                            09-SEP-2002; 2002US-0409293P.
15-JAN-2003; 2003US-0440129P.
20-FEB-2003; 2003WO-US005028
                                                                                                                                                                                                                                                                                                                                        Beigelman L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         51
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADG30034 standard; RNA;
                                                                                                                                                                                                                                                                                                                               ι J, Βε<sub>ιν</sub>
« Usman N,
                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-731676/69.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               of the invention.
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cardiovascular disease; immune disease; rheumatoid arthritis;
                     angiogenic abnormality
                                                                                                                                                                                                                                                                                           WPI; 2004-553738/53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity
Matches 21; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO2004063380-A1
                                                                                      WO2004063380-A1
                                                                                                                                                                                                                                                            Dong X,
                                                                                                                                                                                                                                                                                                                                                                                   abnormalities.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-JUL-2004
                                                                                                                      29-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADQ94243;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                             Series
                                                                                                                                                                                                                                                            Wu Χ,
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IID ADQ9

XX ADQ9

XXX XXX

XXX XX XXX

XXX XXX BS;

KW GPDE

KW GPDE

KW GPDE

KW GAPDE

XX CATOR

XX CATOR

XX CATOR

XX XX ADQ9-29-J
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of the invention demonstrates antiarteriosclerotic, neuroprotective, noctropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary diseases such as restenosis, atherosclerosis, Alzheimer's disease, epilepsy, dementia, huntington's disease, epilepsy, dementia, huntington's disease or amyotrophic lateral sclerosis. Furthermore, the siNA may be utilised for entering the content in the content is that of the siNA RNA RNA RNA RNA ENA content and content in the conten
                                                                                                                                                                                                                                                                                                                                                                                                                                                 New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ss; antiarthritic; antirheumatic; cardiovascular; cytostatic; gene therapy; adeno-associated virus; RNAi; RNA interference; short interfering RNA; gene silencing; short hairpin RNA; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                   Fosnaugh K;
   Alzheimer's, Parkinson's; epilepsy, dementia; huntington's; amyotrophic lateral sclerosis; gene thexapy; ss; hTR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 21; DB 1; Length 21;
Pred. No. 1.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
                                                                                                                                                                                                                                                                                                                                                                   Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 5 A; 4 C; 11 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Short hairpin RNA oligonucleotide SHRNA12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 24; SEQ ID NO 600; 593pp; English.
                                                                                                                                                                                                                                                                                                                                                                   Beigelman L, Chowrira B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GCCTGCCGCCTTCCACCGTTC 156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4.7%; Sc.__
100.0%; Pred
0; N
                                                                                                                                                                                                                                                                                                                                                                                    Thompson J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GCCTGCCGCCTTCCACCGTTC 1
                                                                                                                                                                                                                                                                                                                                  (SIRN-) SIRNA THERAPEUTICS INC
                                                                                                                                                                                                        2002US-0363124P.
2002US-0386782P.
2002US-0406784P.
2002US-0408378P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP
                                                                                                                                                           20-FEB-2003; 2003WO-US005028
                                                                                                                                                                                            2002US-0358580P
                                                                                                                                                                                                                                                                                                  15-JAN-2003; 2003US-0440129P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                    Usman N,
                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-731676/69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                         WO2003074654-A2
                                                                                                                                                                                                                                            29-AUG-2002;
05-SEP-2002;
09-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                   Mcswiggen J,
Jamison S, I
                                                         Unidentified
                                                                                                                                                                                              20-FEB-2002;
                                                                                                                                                                                                              11-MAR-2002;
                                                                                                                                                                                                                              06-JUN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21;
                                                                                                                           12-SEP-2003
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The invention relates to an adeno-associated virus (AAV) that carries a specific RNAi (RNA interference) nucleotide fragment comprising components of the outer shell of a recombinant AAV, a specific RNAi nucleotide fragment, and a control element for regulating the transcription and expression of the RNAi nucleotide fragment carried by encapsulation within the outer shell of a recombinant AAV. The recombinant viruses are useful for inducing RNAi pathway and gene therapy in treating e.g. cancer, cardiovascular diseases, immune diseases like rheumatoid arthritis and angiogenic abnormalities. This sequence corresponds to a short hairpin RNA (shRNA) oligonucleotide used in the
                                                                                                                                                                                                                                                                                                                                                                     recombinant adeno-associated viruses useful for inducing RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                         interference pathway and for gene therapy in treating e.g. cancer, cardiovascular diseases, rheumatoid arthritis and angiogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ss; antiarthritic; antirheumatic; cardiovascular; cytostatic; gene therapy; adeno-associated virus; RNAi; RNA interference; short interfering RNA; gene silencing; short hairpin RNA; cancer; cardiovascular disease; immune disease; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity 100.0%; Fred. No. 1.2
21; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Short hairpin RNA oligonucleotide SHRNA11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 8; SEQ ID NO 20; 53pp; Chinese.
                                                                                                                                                                                                                        Hou Y;
                                                                                                                                             (AGTC-) AGTC GENE TECHNOLOGY CO LTD.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GAAGAGTTGGGCTCTGTCAGC 21
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07-NOV-2003; 2003WO-CN000939.
                                                                      07-NOV-2002; 2002CN-00149319
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Disclosure; Page 24; 79pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-JUL-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27-OCT-1994;
13-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                 inhibition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-JAN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT11035;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 152
원
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                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                         The invention relates to an adeno-associated virus (AAV) that carries a specific RNAi (RNA interference) nucleotide fragment comprising components of the outer shell of a recombinant AAV, a specific RNAi nucleotide fragment, and a control element for regulating the transcription and expression of the RNAi nucleotide fragment carried by encapsulation within the outer shell of a recombinant AAV. The recombinant viruses are useful for inducing RNAi pathway and gene therapy in treating e.g. cancer, cardiovascular diseases, immune diseases like rheumatoid arthritis and angiogenic abnormalities. This sequence corresponds to a short hairpin RNA (shRNA) oligonucleotide used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      rRNA processing; RNA methylation site selection; pseudouridine formation; HACA-enoRNA; telbenerase RNA; cancer; tumour; cell proliferation; trypanosome infection; RNNi-mediated degradation; neo gene; siRNA; small interfering RNA; TER; human; telomerase; RNAi; selection marker.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           snoRNA; small nuclear RNA; box H/ACA; mRNA splicing; mRNA processing;
                                                                                          Series of recombinant adeno-associated viruses useful for inducing RNA interference pathway and for gene therapy in treating e.g. cancer, cardiovascular diseases, rheumatoid arthritis and angiogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New isolated small nuclear RNA (snoRNA) polynucleotides, useful for inducing RNAi-mediated degradation of snoRNA or for treating diseases associated with activity of small nuclear RNA, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                            4.7%; Score 21; DB 1; Length 21;
100.0%; Pred. No. 1.2e+02;
cive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 3 A; 8 C; 3 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    siRNA sequence used for RNA inhibition.
                                                                                                                                                     Example 8; SEQ ID NO 19; 53pp; Chinese.
                      (AGTC-) AGTC GENE TECHNOLOGY CO LTD.
                                                                                                                                                                                                                                                                                                                                                                                             163
                                                                                                                                                                                                                                                                                                                                                                                                                   21
                                                                                                                                                                                                                                                                                                                                                                                                         1 GCCTTCCACCGTTCATTCTAG
                                                                                                                                                                                                                                                                                                                                                                                             143 GCCTTCCACCGTTCATTCTAG
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X,
07-NOV-2002; 2002CN-00149319.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                      21; Conservative
                                             Ma X,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (UYBA-) UNIV BAR-ILAN
                                                                     WPI; 2004-553738/53.
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                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
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                                               איע X, Dong X,
                                                                                                                               abnormalities.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-DEC-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Michaeli S;
                                                                                                                                                                                                                                                                                                  invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADT86999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 151
                                                                                                                                                                                                                                                                                                                                                                        Matches
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that the trypanosome infection is associated with the parasite's snoRNA to that the trypanosome infection is associated with the parasite's snoRNA expression. The polymucleotides of the invention are useful for inducing RNAi-mediated degradation of a small nuclear RNA. They are useful for inducing RNAi-mediated degradation of a small nuclear RNA. They are useful for inducing RNAi-mediated degradation of some polymucleotides of the cancer calls can be inhibited by a polymucleotide sequence which is capable of inducing RNAi mediated degradation of the human telomerase RNA. The snoRNA-2 was found to be part of a gene cluster which includes two additional coding sequences of novel RNA termed as h2 and h3. It has also observed that the snoRNA-2 silencing occurs at the mature RNA transcript level and snoRNA-2 silencing occurs at the snoRNA-2 guided methylation on the S. BrRNA. The expression level of the snoRNA-2 transcripts depend on the orientation of the snoRNA-2 gene with respect to neo gene (selection marker). It has snoRNA-2 gene with respect to neo gene (selection marker). It has snoRNA-2 silencing of sno-RNA-2 is mediated through the uncleus and cytoplasm. The invention suggests that the siRNA mediated snoRNA-2 silencing is not unique to snoRNA cluster-2. It is also snoRNA-2 silencing is not unique to snoRNA cluster-2. It is also complicable for mammalian snoRNAs. The proposed nucleic acid construct comparises of a selection marker gene in reverse orientation. It also includes two promoters and each promoter is capable of directing the
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                                      (small nuclear RNA) molecules or the box H/ACA containing RNA molecules. The method can be used to treat the diseases associated with the activity of small nuclear RNA. Small nuclear RNA molecules are the important regulators of gene expression. They participate in mRNA splicing, mRNA and rRNA processing, RNA methylation site selection and pseudouridine formation (box H/ACA-snoRNA). The telomerase RNA is an important nuclear RNA which serves as a template for telomerase replication. It contains box H/ACA ilke domain which confers the functional localisation of this RNA to the nucleus. The majority of the cancerous tumours contain active
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Telomerase; mammal; antisense; triplex forming oligonucleotide; plasmid;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Antisense oligonucleotide (Tel-AU) inhibiting telomerase activity.
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The present invention provides the method
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94US-00330123.
95US-00387524.
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US09436060A-16.rng.sl

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Villeponteau B,
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                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                              AAT10286;
                                                                                                                                                                                                                                                                                        RESULT 154
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                                                                                                                                            The RNA components of (non) human mammalian telomerase(s) especially from mouse, rat and chinese hamster are all claimed Antisense oligonucleotides can be used to block the activity of the telomerase; probes and primers can be used in detection; vectors and host cells transformed with the isolated telomerase genes can be used for production oligonucleotides and primers was a riborymes and triplex forming claimed can plasmids. A mouse which lacks the telomerase gene therapeutically as can plasmids. A mouse which lacks the telomerase gene the rape and be used for study of telomera regulation in vivo, and the role it plays in immortalisation. The antisense oligonucleotide is synthesised as a 2-0-methyl RNA oligonucleotide and is more resistant to hydrolysis than unmodified RNA oligonucleotides (See AATI1032-35)
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                                                                                                RNA components of (non)human mammalian telomerase(s) - useful in studying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Telomerase; mammal; antisense; triplex forming oligonucleotide; plasmid; probe; primer; ribozyme; ss.
                                                 Marhuenda MA;
                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Antisense oligonucleotide (T3) inhibiting telomerase activity.
                                                                                                                                                                                                                                                                                               4.4%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 1.3e+02; ive 0; Mismatches 0; Indels
                                                 Greider C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Feng J, Funk W, Greider C,
                                                                                                                                                                                                                                                                           Sequence 20 BP; 4 A; 12 C; 3 G; 1 T; 0 U; 0 Other;
                                                Funk W,
                                                                                                          cell senescence and immortalisation.
                                                                                                                           Disclosure; Page 23; 85pp; English.
                                                Feng J,
                                                                                                                                                                                                                                                                                                                                     2 GGTTGCGGAGGGTGGGCCTG 21
                                                                                                                                                                                                                                                                                                                                               20 GGTTGCGGAGGGTGGGCCTG 1
                   SPRING HARBOR LAB
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94US-00330123.
95US-00387524.
95US-00485778.
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95US-00485778
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                                         Villeponteau B;
                                                                                                                                                                                                                                                                                                                  20; Conservative
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                                                                            WPI; 1996-097428/10
                                                                                                                                                                                                                                                                                                        Best Local Similarity
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                           (GERO-) GERON CORP
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07-JUN-1995;
                  (COLD-) COLD
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13-FEB-1995;
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                                                                                                                                                                                                                                                                                               Query Match
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mouse, rat and chinese hamster are all claimed. Antisense of objective to be objectively of the telomerase; probes and primers can be used to block the activity of the telomerase; probes and primers can be used to lock the activity of the telomerase; transformed with the isolated telomerase genes can be used for production of telomerases; RNA and DNA ribozymes and triplex forming oligonucleotides directed against the telomerase genes can be used therapeutically as can plasmids. A mouse which lacks the telomerase gene therapeutically as can plasmids. A mouse which lacks the telomerase gene the role it plays in immortalisation. The antisense oligonucleotide is synthesised as a 2-0-methyl RNA oligonucleotide and is more resistant to hydrolysis than unmodified RNA oligonucleotides (See AATI1032-35)
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RNA components of (non)human mammalian telomerase(s) - useful in studying cell senescence and immortalisation.
                                                                                                                                                                The RNA components of (non) human mammalian telomerase(s) especially from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RNA component of mammalian telomerase, esp. human - useful in identifying e.g. candidate telomerase-modulating agents.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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1.3e+02;
hes 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.4%; Score 20; DB 100.0%; Pred. No. 1.3
                                                                                              Disclosure; Page 23; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 38; 114pp; English.
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94US-00330123.
95US-00472802.
95US-00482115.
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ID AAT10286 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 100.0
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vaccine; cancer; antisense oligonucleotide; ss.
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                                                                                                                                                                                                                                                                                                                                                                                RNA component of mammalian telomerase, esp. human - useful in identifying
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                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence is a RNA component of mammalian telomerase, antisense oligonucleotide, which can be used, along with triple helix forming sequences, to inhibit telomerase activity in cells, esp.
                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Antisense oligonucleotide 14ab for human telomerase RNA component
                                                                                                                                                                             RNA component; mammalian; telomerase; antisense oligonucleotide;
triple helix; inhibition; neoplastic; cells; activity; ss.
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                                                                                                                                                            RNA component of mammalian telomerase antisense oligo Tel-AU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4.4%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 1.3e+02; ive 0; Mismatches 0; Indels
     Length 20;
                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 4 A; 12 C; 3 G; 1 T; 0 U; 0 Other;
     Score 20; DB 1; Le
Pred. No. 1.3e+02;
                                                                                                                                                                                                                                                                                                                                             Andrews WH;
                                                                                                                                                                                                                                                                                                                                                                                        e.g. candidate telomerase-modulating agents
                        Mismatches
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4.4%; Scc...
100.0%; Pre/
0; N
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 GGTTGCGGAGGGTGGCCTG 21
                                                    20 TTTGTCTAACCCTAACTGAG 1
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AAT10289 Btandard; DNA; 20
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Best Local Similarity 100.
Matches 20; Conservative
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                       Conservative
                                                                                                                                                                                                                                                                                                                                                               WPI; 1996-097581/10.
                                                                                                                                                                                                                                                                                                                            (GERO-) GERON CORP.
               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                        neoplastic cells
                                                                                                                                                                                                                                                                                        27-OCT-1994;
07-JUN-1995;
07-JUN-1995;
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   Query Match
Best Local Simi
Matches 20;
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                                                                                                                                          09-SEP-1996
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                                                                                                                                                                                                         Synthetic.
                                                                                                                         AAT10289;
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The present sequence represents an antisense oligonucleotide directed against the human telomerase RNA component gene sequences. The oligonucleotide can be used as an affinity agent in the methods of the invention, which are used to purify human telomerase. The methods involve the use of several sequential steps, including the use of two matrixes that bind molecules bearing negative charges, a matrix that binds molecules bearing negative charges, an affinity purification step and a size separation. Telomerase is a particular target of anticancer therapies, and is useful in assays for characterizing (pre)cancerous cells. Telomerase can also be used to screen for specific modulators, for biochemical analysis of its activity, and in preparation of antibodies. Fragments of telomerase, or nucleic acid encoding them, are used in vaccines, and for treating over expression of telomerase, particularly in
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pruzan RA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Purification of telomerase on affinity material - useful for, e.g diagnosis and treatment of cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Weinrich SL, Atkinson EM, Lichtsteiner SP, Vasserot AP,
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/note= "biotinylated"
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ID AAV41173 Standard; DNA; 20 BP XX
AC AAV41173;
XX
OB-OCT-1998 (first entry)
XX
DT OB-OCT-1998 (first entry)
XX
KN RNA component of human telomerat
XX
KN RNA component; human telomerat
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KW RNA component; human telomerat
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N RNA component
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Matches 20; Conservative
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                                                                                                                                                                     Key
modified_base
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                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                           W09845450-A1
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Synthetic
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WO9828442-A1
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oligonucleotides to the RNA component of human telomerase (hTR). These
antisense oligonucleotides specifically bybridise to a nucleotide
sequence within an accessible region of the hTR, but that does not
bybridise to sequence within the template region of hTR. These
component of human telomerase in a sample. This is useful for diagnosing
cancer (especially neuroblastoma, bladder, colon and prostate cancer),
and providing prognosis for a cancer patient. The inhibitory
cancer (especially neuroblastoma, bladder, colon and prostate cancer),
and providing prognosis for a cancer patient. The inhibitory
cancer (especially neuroblastoma, bladder, colon and prostate cancer),
and providing prognosis for a cancer patient. The inhibitor of
interfering with transcription of the RNA component, decreasing the half-
interfering with transcription of the RNA component, decreasing the half-
cite of the telomerase RNA component transcript, inhibiting assembly of
the RNA component into the telomerase holoenzyme, or inhibiting assembly
colls in vivo. They can be used in therapeutics for treating or
cells in vivo. They can be used in therapeutics for treating or
immunosuppression, and for selectively down-regulating specific branches
cof the immune system, e.g. a specific subset of T-cells, in anti-
inflammatory theraples or for treating infections by, e.g. yeast,
constitutes or fungi
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Wu F, Kealey JT, Pruzan R,
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96US-00770565
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                                                                                                                                                                              19-DEC-1997;
                                                                                                                                                                                                                                                    20-DEC-1996;
20-DEC-1996;
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Sequences shown in AAV41169 to AAV41181 represent antisense
oligonucleotides to the RNA component of human telomerase (hTR). These
antisense oligonucleotides specifically hybridise to a nucleotide
sequence within an accessible region of the hTR, but that does not
hybridise to a sequence within the template region of hTR. These
component of human telomerase in a sample. This is useful for diagnosing
cancer (especially neuroblastoma, bladder, colon and prostate cancer),
and providing prognosis for a cancer patient. The inhibitory
cancer (especially neuroblastoma, bladder, colon and prostate cancer),
and providing prognosis for a cancer patient. The inhibitory
coligonucleotides can inhibit the telomerase activity level in a cell by
interfering with transcription of the RNA component, decreasing the half-
coliforucleotides can inhibiting the telomerase noloenzyme, or inhibiting the be
collymerase activity of telomerase. These antisense oligonucleotides can
cells in vivo. They can be used in therapeutics for treating or
cells in vivo. They can be used in therapeutics for treating or
cells in vivo. and for selectively down-regulating specific branches
confundancery theraptes or for treating infections by, e.g. yeast,
                                                                                                                                                                                                                                                                                                                                                                 - used for detecting
                                                                                                                                                                                                                                                                                                                                                                    New polynucleotide(s) anti:sense to human telomerase - used for detecting or inhibiting human telomerase, e.g. for treating cancers, contraception, immuno-suppression or treating infection.
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                                                                                                                                                                                                                                                       Pruzan R, Weinrich SL;
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                                                      97WO-US023619.
                                                                                                            96US-00770564
                                                                                                                                       96US-00770565
                                                                                                                                                                                                                                                          Kim NW, Wu F, Kealey JT,
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parasites or fungi
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                                                      19-DEC-1997;
                                                                                                                                       20-DEC-1996;
02-JUL-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 159
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04-APR-1997;
02-JUL-1998.
                                                                                                             Kim NW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 161
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Q
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                                                                                                                                                                                                                                               Sequences shown in AAV41169 to AAV41181 represent antisense
oligonucleotides to the RNA component of human telomerase (hTR). These
antisense oligonucleotides specifically hybridise to a nucleotide
equence within an accessible region of the hTR, but that does not
hybridise to a sequence within the template region of hTR. These
component of human telomerase in a sample. This is useful for diagnosing
component of human telomerase in a sample. The inhibitory
and providing prognosis for a cancer patient. The inhibitory
and providing prognosis for a cancer patient. The inhibitory
cligonucleotides can inhibit the telomerase activity level in a cell by
interfering with transcription of the RNA component, decreasing the half-
life of the telomerase RNA component transcript, inhibiting assembly of
the RNA component into the telomerase holoenzyme, or inhibiting the
polymerase activity of telomerase activity in both cultured cells and in
cells in vivo. They can be used in therapeutics for treating or
preventing cancer, for contraception or sterilisation, for
immunosuppression, and for selectively down-regulating specific branches
of the immune system, e.g. a specific subset of T-cells, in anti-
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                                                                                                                                                                        New polynucleotide(s) anti:sense to human telomerase - used for detecting or inhibiting human telomerase, e.g. for treating cancers, contraception, immuno-suppression or treating infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RNA component; human telomerase; antisense oligonucleotide; infection; neuroblastoma; bladder cancer; colon cancer; prostate cancer; cancer; contraception; sterilisation; immunosuppression; therapeutic; hTR; immune system down-regulation; anti-inflammatory therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RNA component of human telomerase (hTR) antisense oligo 20/21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match

4.4%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 1.3e+02;
Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                          Kim NW, Wu F, Kealey JT, Pruzan R, Weinrich SL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 3 A; 7 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             290 CTGCCACCGCGAAGAGTTGG 309
                                                                                                                                                                                                                            Claim 11; Page 65; 80pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20 CTGCCACCGCGAAGAGTTGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAV41180 standard; DNA; 20 BP
                                    97WO-US023619
                                                            96US-00770564
96US-00770565
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                 WPI; 1998-377670/32.
                                                                                                  (GERO-) GERON CORP.
                                    19-DEC-1997;
                                                            20-DEC-1996;
20-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            parasites or
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             02-JUL-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAV41180;
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Sequences shown in AAV41169 to AAV41181 represent antisense cliganuclectides to the RNA component of human telomerase (TTR). These cliganuclectides to the RNA component of human telomerase (TTR). These categorides within an accessible region of the hTR, but that does not hybridise to a sequence within the template region of hTR. These cliganuclectides may specifically be used for detection of an RNA component of human telomerase in a sample. This is useful for diagnosing cancer (especially neuroblastoma, bladder, colon and prostate cancer), and providing prognosis for a cancer patient. The inhibitory coligonuclectides can inhibit the telomerase activity level in a cell by interfering with transcription of the RNA component, decreasing the half-life of the telomerase RNA component transcript, inhibiting assembly of the RNA component into the telomerase. These antisense oligonuclectides can publicate the telomerase activity of telomerase. These antisense oligonuclectides can inhibiting telomerase activity in the collection of the RNA component into the telomerase activity in this collection of the RNA component into the telomerase activity of telomerase activity of telomerase activity of telomerase activity in the collection of the RNA component and in the collection of the RNA component and the RNA c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New polynucleotide(s) anti:sense to human telomerase - used for detecting or inhibiting human telomerase, e.g. for treating cancers, contraception, immuno-suppression or treating infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cells in vivo. They can be used in therapeutics for treating or preventing cancer, for contraception or sterilisation, for immunosuppression, and for selectively down-regulating specific branches of the immune system, e.g. a specific subset of T-cells, in anti-inflammatory therapies or for treating infections by, e.g. yeast, parasites or fungi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Telomerase; human; immune response; cancer; vaccine; treatment; disease;
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                                                                                                                                                                                                                                                                                                                                                                           Weinrich SL;
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                                                                                                                                                                                                                                                                                                                                                                           Wu F, Kealey JT, Pruzan R,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 11; Page 65; 80pp; English.
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97WO-US023619
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96US-00770565
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ses 20; Conservative
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                                                                                                                                                                                                                                                                 (GERO-) GERON CORP.
19-DEC-1997;
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20-DEC-1996;
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associated with overexpression of telomerase e.g. cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 163
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0
                                                                                                                                                                                                                                                                                        This invention describes a novel composition comprising human telomerase having at least 2000-fold (preferably at least 60000-fold) increased relative purity compared with crude extract of cells from adenovirustransformed kidney cell line. The composition is useful for eliciting an immune response in animals and may therefore be used as a vaccine for treating diseases associated with the overexpression of telomerase e.g. cancer. AAZ2105-C22031 represent oligonucleotides used in the isolation of human clone 28-1 which contains a fragment of the human telomerase described in the method of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Compositions comprising human telomerase, useful for treating diseases
                                                                                                                                                                                               treating diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                  Kealey JT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 20; DB 1; Length 20;
Pred. No. 1.3e+02;
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                                                                                                  Pruzan RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                 Compositions comprising human telomerase, useful for treassociated with overexpression of telomerase e.g. cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 0 A; 9 C; 4 G; 7 T; 0 U; 0 Other;
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"5'-biotinylated cytidine"
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                                                                                                  Lichtsteiner SP, Weinrich SL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.4%; Sco...
100.0%; Pred. No....
0; Mismatches
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                                                                                                                                                                                                                                                            Disclosure, Col 45-46; 34pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           361 AGGCCGCAGGAAGACG 380
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                          95US-00510736
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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Best Local Similarity 100.
Matches 20; Conservative
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/note=
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                                                                (GERO-) GERON CORP.
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                            04-AUG-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Atkinson EM,
Vasserot AP;
                                                                                                    Atkinson EM,
Vasserot AP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 162
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The invention relates to promoter regions from mouse and human telomerase RNA (TR) component genes. The TR gene promoter can be linked to a heterologous gene, especially a gene encoding a cytotoxin, for therapy of cancer, especially neoplasias. The telomerase is necessary for the unrestricted proliferative capacity of many human cancers. Mutation or dysregulation of the telomerase expression in cancer. Substances, or upregulation of telomerase expression in cancer. Substances, identified in the methods, can be used to block transcription from the TR gene promoter through interaction of the 5' regulatory sequences. These gubstances, e.g. antisense oligonucleotides, transcription factors, peptide nucleic acids and factors that disrupt signal transduction, are useful for cancer therapy. In particular, gene therapy vectors
                                                              This invention describes a novel composition comprising human telomerase having at least 2000-fold (preferably at least 6000-fold) increased relative purity compared with crude extract of cells from adenovirus-transformed kidney cell line. The composition is useful for eliciting an immune response in animals and may therefore be used as a vaccine for treating diseases associated with the overexpression of telomerase e.g. cancer. AAZ23626-Z23637 represent oligonucleotides used in the isolation of human clone 28-1 which contains a fragment of the human telomerase
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human telomerase RNA gene (hTR) promoter specific primer H11d.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 20; DB 1; Length 20;
Pred. No. 1.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 0 A; 9 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
4.4%; Score 20; DB
Best Local Similarity 100.0%; Pred. No. 1.3
Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                               described in the method of the invention
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Disclosure; Col 49-50; 34pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Fig 12; 109pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       361 AGGCCGCAGGAAGAGGAACG 380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20 AGGCCGCAGGAAGAGGAACG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAZ07301 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-OCT-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1999-479183/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO9938964-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gene therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-AUG-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
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RWA (TR) component genes. The TR gene promoter can be linked to a cancer. Substances, especially a gene encoding a cyctoxin, for therapy of cancer, especially neoplasiase. The telomerase is necessary for the unrestricted proliferative capacity of many human cancers. Mutation or dysregulation of the telomerase repression pathway may cause reactivation or upregulation of telomerase expression in cancer. Substances, cor upregulation of telomerase expression in cancer. Substances, con the methods, can be used to block transcription from the TR gene promoter through interaction of the 5' regulatory sequences. These substances, e.g. antisense oligomuclectides, transcription factors, ceptide nucleic acids and factors that disrupt signal transduction, are useful for cancer therapy. In particular, gene threapy vectors (especially pGT62-codAupp) comprising the promoter and a viral thymidine (substances can be used to convert a prodrug, e.g. gancyclovir, so that neoplasia can be controlled or treated Direct down-regulation of telomerase RNA gene through manipulation of transcription factors may be effective anticancer therapy and the cloning of the hTR gene promoter
                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to promoter regions from mouse and human telomerase
(especially pGT62-codAupp) comprising the promoter and a viral thymidine kinase gene can be used to convert a prodrug, e.g. gancyclovir, so that neoplasia can be controlled or treated. Direct down-regulation of tendencese RNA gene through manipulation of transcription factors may be effective anticancer therapy and the cloning of the hTR gene promoter allows the analysis of therapeutic molecules which modulate hTR promoter activity. Sequences AAA2072696-321 represent PCR primers used in cloning
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             i, TR; promoter; cytotoxin; cancer; neoplasia; hTR;
thymidine kinase gene; anticancer therapy; human;
                                                                                                                                                                                                                                                                                ;
                                                                                                                                                                                                                                  4.4%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 1.38+02; ive 0; Mismatches 0; Indels
                                                                                                                                                   and mutagenisis of human TR gene (hTR) promoter region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human telomerase RNA gene (hTR) specific primer hTRe.
                                                                                                                                                                                          Sequence 20 BP; 1 A; 4 C; 12 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CANC-) CANCER RES CAMPAIGN TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure, Fig 6; 109pp; English.
                                                                                                                                                                                                                                                                                                                                             1 GCCTGGGAGGGGTGGTGGCC 20
                                                                                                                                                                                                                                                                                                                       17 GCCTGGGAGGGGTGGTGGCC 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAZ07275 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99WO-GB000308
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                                                                                                                                                                                                                                                                                20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             relomerase RNA; TR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1999-479183/40.
                                                                                                                                                                                                                                                          Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR primer; 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9938964-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAZ07275;
                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 164
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New peptide nucleic acid (PNA) compounds that inhibit telomerase activity in mammalian cells is useful as probes to detect the RNA component of a mammalian telomerase.
allows the analysis of therapeutic molecules which modulate hTR promoter activity. Sequences AA207623-80 represents PCR primers for amplifying human TR gene (hTR) promoter sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                            Peptide nucleic acid; PNA; telomerase; ribonucleoprotein enzyme; cancer; inhibitor; neoplasia; neurodegenerative disease; aging; hyperplasia; AIDS; HIV; fungal infection; forensic identification; detect; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /*tag= a //note= "Peptide nucleic acid molecule, where N-(2-aminoethyl)glycine units are linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
                                                                                                                                          Gape
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Corey DR;
                                                                                                         Length 20;
                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                               PNA sequence #41 used to inhibit telomerase activity.
                                                                    Sequence 20 BP; 3 A; 5 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                        1.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Shay JW, Norton JC,
                                                                                                       DB 1;
                                                                                                                100.0%; Pred. ....
                                                                                                       Score 20;
Pred. No.
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                                                                                                                                                                             410 CTGAGCTGTGGGACGTGCAC 429
                                                                                                                                                                                                         CTGAGCTGTGGGACGTGCAC 20
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                                                                                                         4.48;
                                                                                                                                                                                                                                                                                                       AAA37583 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Piatyszek MA,
                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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                                                                                                                      Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     paternity testing; ss.
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misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                          AAA37583;
                                                                                                       Query Match
                                                                                                                                                                                                                                                                    RESULT 165
                                                                                                                                                                                                                                                                                      AAA37583,
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diseases characterized by abnormal telomere metabolism or telomerase activity, in combination with antineoplastic and other cyctotoxic or cytostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization probes to detect or quantitate polynucleotides having a human telomerase RNA fhTR) sequence. PNA probes are also used for forensic identification of individuals, e.g. paternity testing, based on hTR gene restriction fragment length polymorphism (RFLP) pattern. PNAs are also useful as probes to detect the RNA component of a mammalian telomerase and as inhibitors of telomerase activity. The method of the present invention allows cancerous conditions to be detected with increased confidence and possibly at an earlier stage, before cells are detected as cancerous based on pathological characteristics. The diagnostic and prognostic methods of the present invention can be used to detect an immortal or neoplastic cell or tumour tissue or cancer of any origin, provided the cell expresses telomerase activity and its RNA component
                                                                                                                                                                                                                                                                                                                                                                                   ;

    .20
    /*tag= a
    /*mote="This sequence is a peptide nucleic acid, i.e. it contains a polyamide backbone instead of a deoxyribose backbone"

                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mammalian; peptide nucleic acid; probe; forensic; paternity testing; human telomerase RNA component; HTR gene RFLP pattern; cancer; inflammation; lymphoproliferative disease; autoimmune disease; neurodegenerative disease; neoplasia; hyperplasia; HTV; AIDS; human immunodeficiency virus; acquired immunodeficiency syndrome; telomere metabolism; mutant; cytostatic; anti-inflammatory; immunosuppressive; polyamide backbone; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer,
                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PNA VIII inhibiting human and mammalian telomerase activity.
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                                                                                                                                                                                                                                                                                                                                              4.4%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 1.3e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 3 A; 5 C; 5 G; 7 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Piatyszek MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                         CTAACCCTAACTGAGAAGGG 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                CTAACCCTAACTGAGAAGGG
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                                                                                                                                                                                                                                                                                                                                                                  1 Similarity 100.
20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Wright WE,
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09-APR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAS15454;
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                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 166
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                                                                                                                                                                                                                                                                                                                                                                                       Matches
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The present invention relates to peptide nucleic acids (PNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in mammalian cells by hybridising to the RNA component of mammalian cells by hybridising to the RNA component of mammalian cells by hybridising to the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polymucleotide having the human telomerase RNA component (hTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal cuspects or unknown descendants based on the hTR gene RFLP pattern. The PNA can be further used for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmacceuticals (such as antineoplastic or cytostatic agents) can be used for treating neoplast, hyperplasia, human immunodeficiency virus (HIV) infections, acquired inmunodeficiency synchome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomera metabolism or telomerase cutivity. The present sequence represents one of the PNA sequences of the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma;
breast; prostate; colon; mixed cell leukaemia; Hodgkin's disease;
fertility; inflammatory condition; tumour; cancer; veterinary;
immunosuppression; telomerase inhibitor; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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inflammation, lymphoproliferative diseases, autoimmune disease,
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                             4.4%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 1.3e+02; cive 0; Mismatches 0; Indels
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/note= "N3'-P5' phosphoramidate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 3 A; 5 C; 5 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human telomerase polynucleotide inhibitor #15.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                    English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         46 CTAACCCTAACTGAGAAGGG 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CTAACCCTAACTGAGAAGGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP
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                  neurodegenerative diseases.
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                                                    29; 46pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-656955/75
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                    Example 1; Col
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modified base
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                                                                                                                                                                                                                                                                                                                                                                                     invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 167
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Matches
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The invention relates to polynucleotide inhibitors (I) and methods for inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity and proliferation of a telomerase positive cell, and in manifacturing a medicament for inhibiting telomerase activity in a cill mand in treating telomerase-mediated condition or disease, such as adenocarcinoma of breast, prostate or colon, mixed cell leukaemia, Hodgkin's disease, fertility and inflammatory conditions. (I) are also useful in treating a tumour or in manufacturing a medicament for the treatment of tumour. The polynucleotide inhibitors may also be used in diagnostic assays for detecting RNA or DNA. Inhibition of telomerase cativity in cells in vivo is useful in prophylactic and therapeutic methods of treating cancer and other disorders involving inappropriate expression of telomerase, and in treating veterinary proliferative for immunosuppression and for selectively down-regulating specific for immunosuppression and for selectively down-regulating specific branches of the immune system. The present sequence represents human telomerase polynucleotide inhibitor #15, as described in the method of
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New polynucleotide useful for inhibiting telomerase activity in cells, for treating telomerase-mediated condition or disease, such as cancers, tumors, Hodgkin's disease, or inflammatory conditions.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Antisense oligonucleotide for human telomerase, Oligo 14ab #1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4.4%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 1.3e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Seguence 20 BP; 3 A; 3 C; 3 G; 11 T; 0 U; 0 Other;
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                                                                                                                                         Claim 8; Page 36; 48pp; English
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97US-00833377.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
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04-APR-1997;
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AASO9477/C
AASO9477/C
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AASO9477/C
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AASO9477/C
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AASO9477/C
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The sequence represents an antisense oligonucleotide used in the purification of human telomerase. The invention relates to a purified human telomerase. The invention relates to a purified human telomerase core enzyme protein comprising 2000-fold increased than the compared with a crude extract of cells from adenovirus-transformed purity compared with a crude activity and a molecular weight of 200-2000 kidney cell line (293 cells) and when associated with telomerase RNA component has DNA polymerase activity and a molecular weight of 200-2000 kilo Daltons (KDa). The purified telomerase is useful for inducing a humoral or cell-mediated immune response in an animal. Purified ciscases associated with over-expression of telomerase, such as cancer and for producing antibodies that recognize telomerase, which are useful as affinity agents in isolating the proteins and for detecting the proteins in a sample, such as cell or tissue. Identification of telomerase aids in diagnosis of cancer or pre-cancerous states.

Telomerase and/or telomerase associated proteins are also useful for screening compounds to identify agents that alter the association of telomerase associated proteins, such as nucleolin or EF2H with telomerase
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Purified human telomerase, useful for inducing immune response in animals, comprises several thousand folds increased purity compared with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pruzan RA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antisense oligonucleotide for human telomerase, Oligo 14ab #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human, Telomerase, vaccine, antibody, cancer; EF2H, nucleolin; antisense oligonucleotide; Oligo 14ab, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Weinrich SL, Atkinson EM, Lichtsteiner SP, Vasserot AP, Kealey \mathtt{JT}_{\mathsf{f}}
                                                                                                                                                                                                                                                                                                                                                          Length 20;
                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 0 A; 9 C; 4 G; 7 T; 0 U; 0 Other;
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100.0%; Pred. No. 1.3e+02;
                                                                                                                                                                                                                                                                                                                                                                    100.0%; Pred. ....
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1es 20; Conservative
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modified_base
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AAS09480/c
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the purification of human telomerase. The invention relates to a purified human telomerase core enzyme protein comprising 2000-fold increased purity compared with a crude extract of cells from ademovinus-transformed kidney cell line (293 cells) and when associated with telomerase RNA component has DNA polymerase activity and a molecular weight of 200-2000 kilo Daltons (KDa). The purified telomerase is useful for inducing a humoral or cell-mediated immune response in an animal. Purified telomerase or immunogenic fragments are useful as vaccines for treating telomerase or immunogenic fragments are useful as vaccines for treating diseases associated with over-expression of telomerase, which are useful as affinity agents in isolating the proteins and for detecting the presence of proteins in a sample, such as cell or tissue. Identification of telomerase and/or telomerase associated proteins are also useful for screening compounds to identify agents that alter the association of telomerase associated proteins are also useful for screening compounds to identify agents that alter the association of telomerase associated proteins are also well for screening compounds to identify agents that alter the association of telomerase.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention provides a method for inhibiting or reducing the growth of a cell which involves administering to the cell a telomere damage inducing agent and a telomerase inhibitory agent. This can be used in the treatment of aberrant cell growth, including cancers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                growth of cell for treating cancer, comprising damage-inducing agent and telomerase inhibitory
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4.4%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 1.3e+02;
Matches 20; Conservative 0; Mismatches 0; Indels
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Pred. No. 1.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
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                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 0 A; 9 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                4.4%; Scc. 100.0%; Pred. No. ...
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         361 AGGCCGCAGGAAGAGAACG 380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AGGCCGCAGGAAGAGGAACG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human telomerase PCR primer #2.
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nes 20; Conservative
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WIENTJES G.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        agent to the cell.
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(WIEN/) WIENT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 170
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This represents an antisense oligonucleotide to the RNA component of human telomerase (hTR). The invention relates to an activator-antisense complex that comprises: (a) an antisense oligonucleotide, complementary comprises: (a) an antisense oligonucleotide, complementary to a 12-25 nucleotide portion of hTR, with a hydroxyl moiety at the first complex and (b) a linker attached to the first end, and (c) an activator of RMsse L attached to the linker. The activator-antisense complex may be used for inhibiting the growth of a telomerase-expressing malignant cells or tumour. The complex is used to specifically cleave the ribonucleotide or tumour, renal cell cancer, mean and prostate cancer. Many other tumour, renal cell cancer, mean and prostate cancer. Many other canner and chronic leukemias, polychemia vera, Hodgkin's and non-Hodgkin's lymphomas, multiple myeloma, Waldenstrom's macroglobulinemia, Hodgkin's lymphomas and solid tumours, including numerous sarcomas and carcinomas. The complex is preferably administered in combination with a chemotherapeutic agent, particularly either displatin, doxornation with a chemotherapeutic agent, particularly either displatin, doxornation with
                                                                                                                                                                                                                                                                    cleave; brain; tumour malignant glioma; breast tumour; renal cell cancer; melanoma; prostate cancer; leukemia; polycthemia vera; myeloma; sarcoma; Hodgkin's lymphoma; Waldenstrom's macroglobulinemia; heavy chain disease; carcinoma; chemotherapeutic; antisense; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mitomycin, daunorubicin, bleomycin, actinomycin D, or neocarzinostatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New RNase L activator-telomerase antisense complex - useful to inhibit
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                                                                                                                                                                                                                                   Human telomerase RNA (hTR) component antisense oligonucleotide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      telomerase activity in telomerase-expressing malignancies
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 19 BP; 6 A; 5 C; 8 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Li G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cowell JK,
20
                         1 GGGTTGCGGAGGGTGGGCCT 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 4; Page 54; 81pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (CLEV-) CLEVELAND CLINIC FOUND. (USSH ) US NAT INST OF HEALTH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     97US-0044507P.
98US-00018125.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98WO-US007397
                                                                                                                           AAV68462 standard; DNA; 19
                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kondo S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1998-609972/51.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
nes 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Silverman RH,
                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-APR-1997;
03-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9847911-A1
                                                                                                                                                                                                  22-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                   AAV68462;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Loca
Matches
                                                                                        RESULT 171
                                                                                                             AAV68462/
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   à
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GTGCTTTTGCTCCCGGGC 94

9/

à

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Gaps

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BP.

AAA37602 standard; RNA; 19

RESULT 173

AAA37602

AAA37602;

15-AUG-2000 (first entry)

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Sequences shown in AAV41169 to AAV41181 represent antisense
cligonucleotides to the RNA component of human telomerase (hTR). These
antisense cligonucleotides specifically hybridise to a nucleotide
sequence within an accessible region of the hTR, but that does not
hybridise to a sequence within the template region of hTR. These
cligonucleotides may specifically be used for detection of an RNA
component of human telomerase in a sample. This is useful for diagnosing
cancer (especially neuroblastoma, bladder, colon and prostate cancer),
and providing prognosis for a cancer patient. The inhibitory
coligonucleotides can inhibit the telomerase activity level in a cell by
interfering with transcription of the RNA component, decreasing the half-
ci life of the telomerase RNA component transcript, inhibiting assembly of
the RNA component into the telomerase holeonizyme, or inhibiting the polymerase activity of telomerase. These antisense oligonucleotides can
be used for inhibiting telomerase activity in both cultured cells and in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New polynucleotide(s) anti:sense to human telomerase - used for detecting or inhibiting human telomerase, e.g. for treating cancers, contraception, immuno-suppression or treating infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cells in vivo. They can be used in therapeutics for treating or preventing cancer, for contraception or sterilisation, for immunosuppression, and for selectively down-regulating specific branches of the immune system, e.g. a specific subset of T-cells, in anti-
                                                                                                                                                                                                                                                                RNA component; human telomerase; antisense oligonucleotide; infection; neuroblastoma; bladder cancer; colon cancer; prostate cancer; cancer; contraception; sterilisation; immunosuppression; therapeutic; hTR; immune system down-regulation; anti-inflammatory therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 inflammatory therapies or for treating infections by, e.g. yeast, parasites or fungi
                                                                                                                                                                                                                         RNA component of human telomerase (hTR) antisense oligo 21ab.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pruzan R, Weinrich SL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 19 BP; 5 A; 3 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 11; Page 65; 80pp; English
19 GTGCTTTTGCTCCCCGCGC 1
                                                                                                    ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   97WO-US023619.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            96US-00770564
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               96US-00770565
                                                                                                    AAV41176 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kim NW, Wu F, Kealey JT,
                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1998-377670/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9828442-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-DEC-1996;
                                                                                                                                                                                   08-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-JUL-1998.
                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                          AAV41176;
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The present sequence represents a telomerase target sequence. The invention relates to an antisense oligonucleotide used as a control sequence alongside a peptide nucleic acid molecule which hybridises to the mammalian telomerase, and inhibits telomerase activity. Telomerase is a ribonucleoprotein enzyme that synthesizes one atraint to the telomeratic DNA, using as a template an il nucleotide contained within the RNA component of the enzyme. The invention capture contained within the RNA encepance of rownponent of the enzyme. The invention contained within the RNA encepance of the sequence of rownponent of the enzyme. The invention contained within the RNA component of the enzyme. The invention contained streams the melting temperature of associating strands, backbone increases the melting temperature of associating strands, controlled spreadation with targeted nucleic acids, and affords greater resistance of degradation by proceases or nucleases. The theorem is neoplasia, hyperplasia, neurodegenerative diseases, aging, human cancers, neoplasia, hyperplasia, neurodegenerative diseases, aging, human immunodeficiency virus (HIV) infection/AIDS (acquired immunodeficiency syndrome) and associated pathologies, fungal infections, and other cytotoxic diseases characterized by abnormal telomerase activity, in combination with antineoplastic and other cytotoxic cytostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled by and seed on the present individuals, e.g. paternity testing, based on hTR gene restriction fragment length polymorphism (RFLP) pattern. PNAs are also useful as probes to detect the RNA component of a mammalian telomerase and as inhibitors of telomerase activity. The method of the present invention allows the control of the present invention allows.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New peptide nucleic acid (PNA) compounds that inhibit telomerase activity in mammalian cells is useful as probes to detect the RNA component of a
                                                                                                                                                             Peptide nucleic acid; PNA; telomerase; ribonucleoprotein enzyme; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 possibly at an earlier stage, before cells are detected as cancerous based on pathological characteristics. The diagnostic and prognostic methods of the present invention can be used to detect an immortal or neoplastic cell or tumour tissue or cancer of any origin, provided the
                                                                                                                                                                                   inhibitor, neoplasia, neurodegenerative disease, aging, hyperplasia, AIDS, HIV, fungal infection, forensic identification, detect; tumour, paternity testing, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Corey DR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              expresses telomerase activity and its RNA component
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Norton JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Shay JW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 2; Col 37; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                   97US-00838545.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              96US-00630019
                                                                                                                  Telomerase target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Piatyszek MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (TEXA ) UNIV TEXAS SYSTEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mammalian telomerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2000-292432/25.
                                                                                                                                                                                                                                                                                                                                                                                                                   09-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                      04-APR-2000.
                                                                                                                                                                                                                                                                                                                           US6046307-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Wright WE,
                                                                                                                                                                                                                                                                             Synthetic.
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Sequence 19 BP; 7 A; 5 C; 3 G; 0 T; 4 U; 0 Other;

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Gaps

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0; Indels

4.2%; Score 19; DB 1; Length 19; 00.0%; Pred. No. 1.5e+02;

100.0%; Prec. ...

19; Conservative

Best Local Similarity

Query Match Matchea

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The present sequence represents an antisense polyamide nucleic acid (PNA)), which is covalently linked to a RNase Lactivator moiety. The oligonuclectide is used in the method of the invention. The specification describes a method of cleaving a specific linvention. The specification describes a method of cleaving a specific RNA, optionally contained within a cell. The method comprises hybridising the RNA with a chimeric molecule of antisense PNA and RNAse L activator molecule is useful for treating Respiratory Syncytial Virus infection, inhibiting tumour growth, treating viral infection (e.g. human immunodeficiency virus (HIV)), autosomal dominant diseases caused by a mutant gene, chronic myelogenous leukemias, and crohns disease and other inflammatory conditions. It is also useful for sequence-specific inhibition of a targeted mRNA in order to effectively knock out the associated gene as an aid to elucidating the function of the encoded
                                  0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cleaving a specific RNA comprising hybridizing the RNA with a chimeric molecule consisting of antisense polyamide nucleic acid (pna) and RNAse L activator moieties, useful for treating e.g. cancer and inflammatory
                                                                                                                                                                                                                                                                                                                                 Antisense polyamide nucleic acid; peptide nucleic acid; PNA; RNase L activator; RNA cleavage; antisense PNA; tumour growth; respiratory syncytial virus infection; viral infection; human immunodeficiency virus; HIV; autosomal dominant disease; chronic myelogenous leukemia; Crohns disease; inflammatory condition; ss.
                                  Gaps
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0
                                                                                                                                                                                                                                                                                                Nucleotide sequence of an antisense polyamide oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Van Der Marel GA;
Length 19;
                                  0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19 BP; 6 A; 5 C; 8 G; 0 T; 0 U; 0 Other;
Score 19; DB 1; Pred. No. 1.5e+02; 4; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Verheijen JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US DEPT HEALTH & HUMAN SERVICES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 25; Page 32; 41pp; English.
                                                                       62
                                                                                                        cucua acccua a cuca ca a 19
                                                                                                                                                                                                 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      98US-0099173P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99WO-US020159.
                                                                         GTCTAACCCTAACTGAGAA
   4.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Van Boom JH,
                                                                                                                                                                                                 AAZ99775 standard; DNA; 19
                                                                                                                                                                                                                                                                   (first entry)
   Query Match
Best Local Similarity 78.9
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (UYLE-) RIJKSUNIV LEIDEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-256971/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200014219-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        04-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Torrence PF,
                                                                                                                                                                                                                                                                     12-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16-MAR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                  AAZ99775;
                                                                           44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (USSH)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     protein
                                                                                                                                                              RESULT 174
                                                                                                                                                                                  AAZ99775
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The present invention relates to peptide nucleic acids (PNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in nammalian cells by Mybridising to the RNA component of mammalian cells by Mybridising to the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polymucleotide having the human telomerase RNA component (hTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the hTR gene RFLP patern. The PNAs can be further used for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases, the PNAs in combination with other pharmaceuticals (such as antineoplastic or cytostatic agents) can be used for treating neoplasia, hyperplasia, human immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomerase metabolism or telomerase activity. The present sequence repersenting hTR RNA strand #3 is used to test the inhibition of telomerase activity by the PNAs of the present
                                                                                                                                                                                                                                                         Mammalian, forensic, paternity testing, human telomerase RNA component, hTR gene RFLP pattern, cancer, inflammation; lymphoproliferative disease; autofimmune disease, neurodegenerative disease, neoplasia; hyperplasia; HIV, AIDS; human immunodeficiency virus; telomere metabolism; cytostatic; acquired immunodeficiency syndrome; anti-inflammatory; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Norton JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 19; DB 1; Length 19;
Pred. No. 1.5e+02;
4; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 19 BP; 7 A; 5 C; 3 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Piatyszek MA, Corey DR,
                                                                                                                                                                                                                           Human telomerase RNA (hTR) strand #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 2; Col 37-38; 46pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 GUCUAACCCUAACUGAGAA 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GICTAACCCTAACTGAGAA 62
                 19 Grecrrrrecrecedes 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99US-00349532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  96US-00630019.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  97US-00838545
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4.28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 Similarity 78.9%;
15; Conservative
neurodegenerative diseases
                                                                                                              AAS15473 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (TEXA ) UNIV TEXAS SYSTEM.
                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wright WE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-638024/73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                     US6294650-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             08-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-APR-1997;
                                                                                                                                                                                        14-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-SEP-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nvention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    44
                                                                                                                                                      AAS15473;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Shay JW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Loca
Matches
                                                                           RESULT 175
                                                                                                AAS15473
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Gaps

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4.2%; Score 19; DB 1; Length 19; 100.0%; Pred. No. 1.5e+02; ive 0; Mismatches 0; Indels

Query Match Best Local Similarity 100.0 Matches 19; Conservative

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Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micoro-RNA; miRNA; short historing RNA; siRNA; double-stranded RNA; micoro-RNA; miRNA; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and genemapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human TERC-targeted double-stranded siNA, which is identical to the c-fos transcript target sequence.
                                                                                             Human TERC transcript target sequence/siNA upper strand, SEQ ID 20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 3; SEQ ID NO 20; 145pp; English.
                       ADF93303 standard; RNA; 19 BP
                                                                                                                                                                                                                                                                                                                              2002US-0358580P
2002US-0363124P
                                                                                                                                                                                                                                                                                                                                                    2002US-0386782P.
2002US-0396600P.
2002US-0406784P.
                                                                                                                                                                                                                                                                                                                                                                                                     09-SEP-2002; 2002US-0409293P.
15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                                                                                                       11-FEB-2003; 2003WO-US004088
                                                                                                                                                                                                                                                                                                                                                                                          2002US-0408378P
                                                                                                                                                                                                                                                                                                                                                                                                                                        (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mcswiggen J, Beigelman L;
                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-689777/65.
                                                                                                                                                                                                                                                       WO2003070742-A1.
                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                              20-FEB-2002;
                                                                                                                                                                                                                                                                                                                                                     06-JUN-2002;
                                                                                                                                                                                                                                                                                                                                                                   17-JUL-2002;
                                                                                                                                                                                                                                                                                                                                                                               29-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                                         05-SEP-2002;
                                                                                                                                                                                                                                                                                                                                           11-MAR-2002;
                                                                      26-FEB-2004
                                                                                                                                                                                                                                                                                28-AUG-2003.
                                               ADF93303;
RESULT 176
            ADF93303
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and
                                                                                                                                                                                                                                                                                                                                                                                                                        Cytostatic, vasotropic, protozoacide, immunosuppressive, dermatological, neuroprotective, anti-HIV, ophthalmological; antiulcer, antirheumatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                               antiarthritic; antiinflammatory, gene therapy, telomerase; human; terc; RNA interference; short interfering nucleic acid; slNA; short interfering RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                    Human TERC transcript target sequence/siNA upper strand, SEQ ID 24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
                                                                                          ;
0
                                                Query Match 4.2%; Score 19; DB 1; Length 19; Best Local Similarity 89.5%; Pred. No. 1.5e+02; Matches 17; Conservative 2; Mismatches 0; Indels
              Sequence 19 BP; 1 A; 7 C; 9 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 3; SEQ ID NO 24; 145pp; English.
                                                                                                                                248 CTGGAGGCCGCGGTCGGCC 266
                                                                                                                                                                       1 cudandecedecedece 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2002US-0358580P.
2002US-0363124P.
2002US-0386782P.
2002US-0406784P.
2002US-0408784P.
2002US-0408378P.
                                                                                                                                                                                                                                                                   ВР
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                                                                                                                                                                                                                                                                   ADF93307 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mcswiggen J, Beigelman L;
                                                                                                                                                                                                                                                                                                                                              26-FEB-2004 (first entry)
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05-SEP-2002;
09-SEP-2002;
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11-MAR-2002;
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US09436060A-16.rng.sl

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 used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human TERC-targeted double-stranded siNA, which is identical to the c-fos transcript target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; siNA; oble-stranded RNA; mirro-RNA; miRNA; short hairpin RNA; siRNA; compression modulation; gene therapy; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
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                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human TERC transcript target sequence/siNA upper strand, SEQ ID 29.
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                                                                                                                                                                                                        Sequence 19 BP; 0 A; 7 C; 9 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 3; SEQ ID NO 29; 145pp; English.
                                                                                                                                                                                                                                                                                                                      320 CCGCGGGTCTCTCGGGGGC 338
                                                                                                                                                                                                                                                                                                                                            1 CCGCGGGUCUCGGGGGGC 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        11-MAR-2002; 2002US-0363124P.
06-UUN-2002; 2002US-0366782P.
17-UUL-2002; 2002US-039660P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADF93312 standard; RNA; 19 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                             Local Similarity 84.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-689777/65.
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Matches
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cc regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short contrained that sinAs include short hairpin RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short chairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised.

CC expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are consed to omboliate expression of the telomerase genes in cells, tissue consed to omboliate expression of the telomerase genes in cells, tissue consed for treating cancer, restencist, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, consed for treating cancer, restencist, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, consed for understand referencies, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene corresponded to the correspondence of a human TERC-teargeted double-stranded correspondence in Na which is identical to the correspondence of a human TERC-teargeted double-stranded
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4.2%; Score 19; DB 1; Length 19; 78.9%; Pred. No. 1.5e+02; trive 4; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           410 CTGAGCTGTGGGACGTGCA 428
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2002US-0396600P.
2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
2003US-0440129P.
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2002US-0363124P.
2002US-0386782P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 78.5
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29-AUG-2002; 2
05-SEP-2002; 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mcswiggen J,
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WPI; 2003-689777/65.

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA connergulate expression of the one or more telomerase genes by RNA connergulate expression of may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense could be stranded. They further comprise sense and antisense con antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain a vertor or enzymatically synthesised. The invention also capressed from a vector or enzymatically synthesised. The invention also captes to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are captained or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be conditions to the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, lupus erythematosus, ALDS, macular degeneration, and gene constition and validation, genetic engineering, pharmacogenomics, studying gene function, and gene constition and sent sequence constition and sent sequ
                               New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
                                                                                                                                          Example 3; SEQ ID NO 287; 145pp; English.
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4.2%; Score 19; DB 1; Length 19; 100.0%; Pred. No. 1.5e+02; ive 0; Mismatches 0; Indels Sequence 19 BP; 5 A; 7 C; 4 G; 0 T; 3 U; 0 Other; Query Match

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Gaps

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302 AGAGTIGGGCTCTGTCAGC 320 19 AGAGTTGGGCTCTGTCAGC 1 19; Conservative Best Local Similarity Matches 19; Conserv g ઠે

ADF93564 standard; RNA; 19 BP 26-FEB-2004 (first entry) ADF93564; RESULT 180 ADF93564/c

Human TERC siNA lower strand, SEQ ID 291.

Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short histopin RNA; shraw, expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.

Homo sapiens.

WO2003070742-A1.

28-AUG-2003

11-FEB-2003; 2003WO-US004088

20-FEB-2002;

2002US-0363124P. 2002US-0386782P. 2002US-0396600P. 11-MAR-2002; 06-JUN-2002; 17-JUL-2002;

The invention relates to short interfering nucleic acids (sinA) which downregulate expression of the one or more telomerase genes by RNA interference. The sinAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antiense regions, or alternatively are assembled from a sense oligonucleotide and an arisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telemerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematchosus, AIDS, macular degeneration, skin ulcers and rheumatchid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, suddying gene function, and gene mapping (e.g., of single nuclecide polymorphisms). The present sequence represents the lower strand of a human TERC-targeted double-stranded New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer. Example 3; SEQ ID NO 291; 145pp; English. 29-AUG-2002; 2002US-0406784P. 05-SBP-2002; 2002US-0408378P. 09-SEP-2002; 2002US-0409293P. 15-JAN-2003; 2003US-0440129P. (RIBO-) RIBOZYME PHARM INC. Beigelman L; WPI; 2003-689777/65. Mcswiggen J,

Sequence 19 BP; 1 A; 7 C; 6 G; 0 T; 5 U; 0 Other;

Gaps ö Length 19; 0; Indels Query Match
4.2%; Score 19; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 1.5e+02;
Matches 19; Conservative 0; Mismatches 0;

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374 AGGAACGGAGCGAGTCCCC 392 19 AGGAACGGAGCGAGTCCCC 1 RESULT 181 g ठ

(first entry) 26-FEB-2004 ADF93294;

ADF93294 standard; RNA; 19 BP

Human TERC transcript target sequence/siNA upper strand, SEQ ID 11.

Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short halfpin RNA; shrNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.

Homo sapiens

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neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic;
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Matches
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                                                                                                                                                                                                                                                                                                                                                    The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA commerced interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense of digonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue capinates for the treatment of a variety of conditions. They may be transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, informance or age-related diseases, e.g. multiple sclerosis, lugues erythematosus, AIDS, macular degeneration, sin dagnosis, therapeutic target identification and validation, shin ulcers and rheumatoid arthritis. The siNAs are also useful for drug capine: the upper strand of a human TERC-targeted double-stranded capines tranded of a human TERC-targeted double-stranded condition, which is identical to the c-fos transcript target sequence.
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                                                                                                                                                                                                                                                                                          short interfering nucleic acid downregulates expression of the
                                                                                                                                                                                                                                                                                                       telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                                                                                                                                                                                                                                                 Example 3; SEQ ID NO 11; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          86 TCCCCGCGCGCTGTTTTC 104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-MAR-2002; 2002US-0363124P.
06-JUN-2002; 2002US-0366782P.
17-JUL-2002; 2002US-0396600P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0409239P.
15-JAN-2003; 2003US-040129P.
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                                                       11-FEB-2003; 2003WO-US004088
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Best Local Similarity 63.2;
....heg 12; Conservative
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  WO2003070742-A1
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                            28-AUG-2003
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The invention relates to short interfering nucleic acids (siNh) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double stranded RNA, micro-RNA (miRNA) and short can contain deoxyribonucleotides, and can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised.

CC expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vitro of in vitro of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in calls, tissue cyprosace to conganisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AlDS, macular degeneration, skin ulcers and rheumatoid architis. The sinks are also useful for drug coreening, diagnosis, lupus erythematosus, AlDS, macular degeneration, careening, diagnosis, laberatic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence consents the upper strand of a human TERC-targeted double-stranded sina, which is identical to the represent strand of a human TERC-targeted double-stranded sina.
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antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ;
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Pred. No. 1.5e+02;
5; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 19 BP; 2 A; 6 C; 6 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 3; SEQ ID NO 16; 145pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1:||||:||||:
1 AUGUCAGCUGCUGGCCCGU 19
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2002US-0386782P.
2002US-0396600P.
2002US-0406784P.
2002US-0408378P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11-FEB-2003; 2003WO-US004088.
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Local Similarity 73.7%;
es 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                     WO2003070742-A1.
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05-SEP-2002;
09-SEP-2002;
                                                                                                                                                                                                                                                                                                          Homo sapiens.
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06-JUN-2002;
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Length 19;

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Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; slNA; short interfering nucleic acid; slNA; short interfering RNA; shRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                               Human TERC transcript target sequence/sinA upper strand, SEQ ID 17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
                                                                                                                                                                                                                                                                                                               11-MAR-2002; 2002US-0363124P.

06-UTN-2002; 2002US-03867B2P.

17-UTL-2002; 2002US-0396600P.

29-AUG-2002; 2002US-0406784P.

05-SSP-2002; 2002US-0408378P.

15-JAN-2003; 2003US-040933P.
            ADF93300 standard; RNA; 19 BP
                                                                                                                                                                                                                                                                                11-FEB-2003; 2003WO-US004088
                                                                                                                                                                                                                                                                                                                                                                                                            (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                   Beigelman L;
                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-689777/65.
                                                                                                                                                                                                                                  WO2003070742-A1.
                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                   Mcswiggen J,
                                                                                                                                                                                                                                                                                                       20-FEB-2002;
                                                         26-FEB-2004
                                                                                                                                                                                                                                                         28-AUG-2003
                                  ADF93300;
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonuclectide and an interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised, captessed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue captants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus exthematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human TERC-targeted double-stranded which is identical to the c-fos transcript target sequence.

Example 3; SEQ ID NO 17; 145pp; English.

Seguence 19 BP; 1 A; 10 C; 5 G; 0 T; 3 U; 0 Other;

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA controlled and and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised. Expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue used to modulate expression of the telomerase genes in cells, tissue transplants or roganisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically
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                                                                                                                                                                                                                                                                                                                                                                                Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological;
                                                                                                                                                                                                                                                                                                                                                                                              neuroprotective, anti-HIV; ophthalmological; antillicer; antitheumatic; antiantucer; antitheumatic; antiantucer; antitheumatic; antiantucer; antitheumatic; antiantucer; antitheumatic; antiantucer; and therefore a cid; alwa; terc; RNA interference; short interfering nucleic acid; siNA; alkNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosia; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protozoal), transplant rejection, or autoimmune or age-related diseases
                                           Gaps
                                                                                                                                                                                                                                                                                                                                          Human TERC transcript target sequence/siNA upper strand, SEQ ID 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
                                           ö
Score 19; DB 1; Length 19;
Pred. No. 1.5e+02;
3; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 3; SEQ ID NO 23; 145pp; English.
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                                                                               194 TTCGCCCTCCCGGGGACC 212
                                                                                                                    1 uncececececeses 19
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; 2002US-0386782P.
; 2002US-039660P.
; 2002US-0406784P.
; 2002US-0408378P.
                                                                                                                                                                                                                      ВР
  4.2%;
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                                                                                                                                                                                                                      ADF93306 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                     26-FEB-2004 (first entry)
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                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-689777/65.
  Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO2003070742-A1.
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                                                                                                                                                                                                                                                             ADF93306;
                                                                                                                                                                                  RESULT 184
                                                                                                                                                                                                   ADF93306
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US09436060A-16.rng.sl

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiartritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short interfering syr, ene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
e.g. multiple sclerosis, lupus erytnematosus, Allo, Machine construction and rheumatoid arthritis. The slMAs are also useful for drug screening, diagnosis, therapeutic taget identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human TERC-targeted double-stranded siNA, which is identical to the c-fos transcript target sequence.
                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human TERC transcript target sequence/siNA upper strand, SEQ ID 31,
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                                                                                                                                                                                     Score 19; DB 1; Length 19; Pred. No. 1.5e+02;
                                                                                                                                                                                                                          0; Indels
                                                                                                                                                Sequence 19 BP; 3 A; 4 C; 7 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                          5; Mismatches
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                                                                                                                                                                                                                                                               302 AGAGTTGGGCTCTGTCAGC 320
                                                                                                                                                                                                                                                                                    AGAGUUGGGCUCUGUCAGC 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-FEB-2002; 2002US-0358580P.
11-MAR-2002; 2002US-0353124P.
06-JUN-2002; 2002US-0386782P.
17-JUL-2002; 2002US-0396600P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408738P.
                                                                                                                                                                                                                                                                                                                                                                                             ADF93314 standard; RNA; 19 BP
                                                                                                                                                                                 4.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                            14; Conservative
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                                                                                                                                                                                     Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-FEB-2004
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expressed from a vector or enzymatically synthesised, relates to kits for the in vivo delivery of sink, conjugates and/or complexes of sink, and vectors that expressed. The invention also relates to kits for the in vivo or in vivo delivery of sink, conjugates and/or complexes of sink, and vectors that express sink. The sinks are used to modulate expression of the telomerase genes in cells, tissue explaints or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, c.g. multiple sclerosis, lupus erythematosus, ALDS, macular degeneration, skin ulcers and rheumatoid arthritis. The sinks are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence mapping (e.g., of single nucleotide polymorphisms). The present sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antianfilammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering nucleic acid; siNA; mirro-RNA; mirro-RNA; shrNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified,
                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
                                                                                                                                                                                                                                                                                                                     siNA, which is identical to the c-fos transcript target sequence
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0
                                                                                                                                                                                                                                                                                                                                                                                              4.2%; Score 19; DB 1; Length 19; 84.2%; Pred. No. 1.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                       Sequence 19 BP; 5 A; 7 C; 4 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                     3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human TERC sinA lower strand, SEQ ID 290.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP.
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2002US-0386782P.
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2002US-0406784P.
2002US-0408378P.
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11-MAR-2002; 2
06-JUN-2002; 2
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29-AUG-2002;
05-SEP-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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15-JAN-2003; 2003US-0440129P.

(RIBO-) RIBOZYME PHARM INC.

dcswiggen J, Beigelman L;

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised. expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express in cells, tissue cused to modulate expression of the telomerase genes in cells, tissue cused to modulate expression of the telomerase genes in cells, tissue cused to recaming (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisme). The present sequence represents the lower strand of a human TERC-targeted double-stranded
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                                   Example 3; SEQ ID NO 290; 145pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CTTTCAGGCCGCAGGAGA 1
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2002US-0386782P.
2002US-0396600P.
2002US-0406784P.
2002US-0408378P.
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05-SEP-2002;
09-SEP-2002;
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06-JUN-2002;
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise sense and antisense of interference. The siNAs may or may not comprise sense and antisense regions, or alternatively are assembled from a sense onigonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short contrain decayribonucleotides, and can be unmodified or chemically modified. Can contain decayribonucleotides, and can be chemically synthesised, can contain decayribonucleotides, and can be chemically synthesised, can contain decayribonucleotides, and can be chemically synthesised. Can complexes of siNA; and vectors that express fine invention also relates to kits for the in vitro or in vitro delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treatment or a variety of conditions. They may be used for treatment rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug sortenting, diagnosis, lharapeutic carget identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence corpused the capared and contains and the capared and contains and sense of a human TRRC-targeted double-stranded and sequence and which is identical to the capared and contains and sense and some contains and sense and sequence and sequence mapping (e.g., of single nucleotide polymorphisms). The present sequence and success the contains and sense and some contains and sense and some contains and sense and some contains and sense and contains and sense contains and sense contains and sense co
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                                                                                                                                                                                                          New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 19 BP; 1 A; 7 C; 8 G; 0 T; 3 U; 0 Other;
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                                                                                                                                                                                                                                                                                          Example 3; SEQ ID NO 21; 145pp; English.
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Best Local Similarity 84.2 Matches 16, Conservative
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonuclectide and an trisense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short harpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically modified, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates cand/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically contozoal), transplant rejection, or autoimmune or age-related diseases, e.g. e.g. multiple sclerosis, lubus erythematosus, ADS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a human TERC-targeted double-stranded
                                                                                                                                                                                                                                                                                                                                                                                  New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                   2002US-0386782P.
2002US-0396600P.
2002US-0406784P.
2002US-0408378P.
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                                                                                 2002US-0363124P
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                  11-FEB-2003; 2003WO-US004088
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05-SEP-2002;
09-SEP-2002;
                                                                                                   06-JUN-2002;
17-JUL-2002;
                                                            20-FEB-2002;
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense or regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short can contain deoxyribonucleotides, and can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised. The invention also relates to kits for the in vitro of in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for trearging cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases. Sin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, lupus erythematosus, AIDS, macular degeneration, genetic engineering, pharmacogenomics, studying gene function, and gene cappands (e.g., of single nucleotide polymorphisms). The present sequence mapping (e.g., of single nucleotide polymorphisms). The present sequence the single of the single cargeded cappands of the single cappends of the single cappered cappands of the single cappered cappands of the single polymorphisms). The present sequence cappands of the single cappands of t
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                       short hairpin RNA; shRNA; expression modulation; gene therapy;
drug screening; diagnosis; therapeutic target identification;
pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
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short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA;
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100.0%; Pred. No. 1.5e+02;
.ive 0; Mismatches 0; Indels
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11-WAR-2002; 2002US-0361124P.

06-UUN-2002; 2002US-0386782P.

17-UUL-2002; 2002US-0396600P.

29-AUG-2002; 2002US-0406784P.

09-SEP-2002; 2002US-0408318P.

09-SEP-2002; 2002US-0408318P.

15-JAN-2003; 2003US-0440129P.
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ID ADF93301 standard; RNA; 19 BP.
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                                                                                                                               Homo sapiens.
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA comparisons. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense coligonucleotide. Specifically, the siNAs include short interfering RNA (shRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised. Complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in calls, tissue candor complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in calls, tissue captants or organisms (e.g., by ex vivo gene therapy), or in grafts and cransplants for the treatment of a variety of conditions. They may be used for treatming cancer, restenosis, induces genes in calls, degeneration, processed in ulcers and rheumatoid arthritis. These macular degeneration, skin ulcers and rheumatoid arthritis. The present sequence captenesing, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human TERC-target educated considering.
                                                                                                            Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; mioro-RNA; mixAd; short hairpin RNA; shraw, expression medulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
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                                                                              Human TERC transcript target sequence/siNA upper strand, SEQ ID 18
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05-SEP-2002, 2002US-0408378P-
09-SEP-2002, 2002US-04902939P-
15-JAN-2003, 2003US-0440129P-
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2002US-0386782P.
2002US-0396600P.
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                                      (first entry)
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                                      26-FEB-2004
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  ADF93301;
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                                 4.2%; Score 19; DB 1; Length 19; 84.2%; Pred. No. 1.56+02; ive 3; Mismatches 0; Indels
Sequence 19 BP; 0 A; 8 C; 8 G; 0 T; 3 U; 0 Other;
                                                                        16; Conservative
                                                      Best Local Similarity
                                   Query Match
                                                                      Matches
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The invention relates to short interfering nucleic acids (sINA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised. Can contain deoxyribonucleotides, and can be chemically synthesised. Can complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue cand/or complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue cand/or complexes of siNA, and vectors that express siNA. The siNAs are used to roganisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically concount or and transplant rejection, or autoimmune or age-related diseases. Can multiple sclerosis, lupus erythematesus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation,
                                                                                                                                                                                                                                                                                                                          Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering uncleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; mioro-RNA; minn, short interfering RNA; siRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
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212 CTGCGGCGGGTCGCCTGCC 230
                    ADF93553 standard; RNA; 19 BP
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2002US-0408378P.
2002US-0409293P.
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2002US-0386782P.
2002US-0396600P.
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06-JUN-2002;
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genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a human TERC-targeted double-stranded
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                                                                                                                      Sequence 19 BP; 5 A; 6 C; 6 G; 0 T; 2 U; 0 Other;
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                                                                                                                                                                                                                                                                     ATGTCAGCTGCTGGCCCGT 194
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06-UUN-2002; 2002US-0386782P.
17-UUL-2002; 2002US-036600P.
29-AUG-2002; 2002US-0406784P.
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2003US-0440129P.
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Best Local Similarity 100.
Matches 19; Conservative
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                   and/or complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therappy, or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoinmune or age-related diseases, e.g. multiple sclerosis, lupus expthematosus, AIDS, macular degeneration skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a human TERC-targeted double-stranded
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relates to kits for the in vitro or in vivo delivery of siNA; conjugates
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100.0%; Pred. No. 1.5e+02;
tive 0; Mismatches 0; Indels
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06-JUN-2002; 2002US-0366782P.
17-JUL-2002; 2002US-0396600P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2003US-040923P.
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Mcswiggen J,

WPI; 2003-689777/65.

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CC The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (shRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (shRNA). The siNAs can be unmodified or chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. Complexes of siNA; and vectors that express since link, inconjugates and/or complexes of siNA; and vectors that express in cells, tissue used to modulate expression of the telomerase genes in cells, tissue cused to modulate expression of the telomerase genes in cells, tissue cused to modulate spression of the telomerase genes in cells, tissue cused to modulate spression of the vivo gene therapy), or in grafts and creating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, cused for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases. Cused for ulcers and rheumatoid arthritis. The sinAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, shin ulcers and rheumatoid arthritis. The sinAs are also useful for drug screening, diagnosis, therapeutic playing gene function and validation, wapping (e.g., of single nucleotide polymorphisms). The present sequence ceptores tepresents the upper strand of a human TERC-targeted double-stranded cusples and related to the c-fos transcript target sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4.2%; Score 19; DB 1; Length 19; 94.7%; Pred. No. 1.5e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 19 BP; 5 A; 10 C; 3 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      284 CACCCACTGCCACCGCGAA 302
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2002US-0408378P.
2002US-0409293P.
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2002US-0386782P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2002US-0358580P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity 94.7
nes 18; Conservative
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06-JUN-2002; 2
17-JUL-2002; 2
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05-SEP-2002; 2
09-SEP-2002; 2
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2003US-0440129P

15-JAN-2003;

(RIBO-) RIBOZYME PHARM INC.

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise shonclectides and may be double or single stranded. They further comprise sense and mathense regions, or alternatively are assembled from a sense oligonuclectide and an artisense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised. Can contain deoxyribonuclectides, and can be chemically sinhesised, can contain a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are cused to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular despenaration, skin ulcers and rheumatoid arthritis. The siMAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, appending engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a human TERC-targeted double-stranded
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                   New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19 BP; 7 A; 6 C; 5 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                            Example 3; SEQ ID NO 274; 145pp; English.
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Matches
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pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to short interfering nucleic acids (siNh) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonuclectide and an antisense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue caplants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapebutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence mapping (e.g., of single nucleotide polymorphisms). The present sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering mucleic acid; slNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; micro-RNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification;
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100.0%; Pred. No. 1.5e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 19 BP; 7 A; 4 C; 8 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 3; SEQ ID NO 275; 145pp; English.
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                                                        2002US-0396782P.
2002US-0396600P.
2002US-0406784P.
                                                                                                                             2002US-0408378P
2002US-0409293P
                                                                                                                                                                                 15-JAN-2003; 2003US-0440129P
            2002US-0358580P
                                 2002US-0363124P
                                                                                                                                                                                                                                  (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                Beigelman L;
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Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                 WPI; 2003-689777/65.
                                                                               17-JUL-2002;
29-AUG-2002;
05-SEP-2002;
09-SEP-2002;
       20-FEB-2002;
11-MAR-2002;
06-JUN-2002;
                                                                                                                                                                                                                                                                                     Mcswiggen J,
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonuclectide and an antisense oligonuclectide. Specifically, the siNAs include short hairpin RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised. Septement of the invention also relates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue transplants for the treatment of a variety of conditions. They may be transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infections diseases (specifically conditions).
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ID ADF93555 standard; RNA; 19 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                    29-AUG-2002; 2002US-0406784P
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                                                                                                                                                          06-JUN-2002; 2002US-0386782P.
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                                                                                                                                                                                                                                                                                                                                  2002US-0363124P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-689777/65.
                                                                  WO2003070742-A1
Homo sapiens.
                                                                                                                                                                                                                                                                                                                               11-MAR-2002;
06-JUN-2002;
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Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological;
                                          neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiathritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short haltpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                                                                                                                                                                                                                                              New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 3; SEQ ID NO 282; 145pp; English.
          Human TERC siNA lower strand, SEQ ID 282
                                                                                                                                                                                                                                                                              29-AUG-2002; 2002US-0406784P.
0S-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
15-JAN-2003; 2003US-0440129P.
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2002US-0386782P.
2002US-0396600P.
                                                                                                                                                                                                            11-FEB-2003; 2003WO-US004088
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                                                                                                                                                                                                                                                                                                                                                                 Beigelman L;
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                                                                                                                                                             WO2003070742-A1.
                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                  Mcswiggen J,
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06-JUN-2002;
                                                                                                                                                                                                                                    20-PEB-2002;
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense or regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double stranded RNA, micro-RNA (miRNA) and short can contain deoxyribonucleotides, and can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can complexes of siNA; and vectors that express sinh. The siNAs are used to modulate expression of the telomerase genes in cells, tissue cexplants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically proteozoal), transplant rejection, or autoimmune or age-related diseases, es multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug certening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence contains the lower strand of a human TERC-targeted double-stranded

Seguence 19 BP; 3 A; 8 C; 8 G; 0 T; 0 U; 0 Other;

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                      Gaps
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0
4.2%; Score 19; DB 1; Length 19; 00.0%; Pred. No. 1.5e+02;
                      0; Indels
       100.0%; Prec. ...
                       19; Conservative
Query Match
Best Local Similarity
                        Matches
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19 CTGCGGCGGGTCGCCTGCC 1

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ADF93556 standard; RNA; 19 BP
   26-FEB-2004 (first entry)
  ADF93556;
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Human TERC siNA lower strand, SEQ ID 283

Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; mixno-RNA; abort hairpin RNA; shawa; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.

Homo sapiens.

WO2003070742-A1.

28-AUG-2003

2002US-0363124P. 2002US-0386782P. 2002US-0396600P. 2002US-0406784P. 2002US-0408378P. 2002US-0409293P. 11-FEB-2003; 2003WO-US004088 2002US-0358580P 29-AUG-2002; 05-SEP-2002; 09-SEP-2002; 11-MAR-2002; 06-JUN-2002; 20-FEB-2002; 17-JUL-2002;

2003US-0440129P (RIBO-) RIBOZYME PHARM INC. LS-JAN-2003;

Mcswiggen J, Beigelman L;

WPI; 2003-689777/65.

New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.

Example 3; SEQ ID NO 283; 145pp; English.

The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonuclectide and an interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised. Concaptessed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA, and vectors that express siNA. The siNAs are used to modilate expression of the telahomerase genes in cells, issue to replace to respension of the telahomerase genes in cells, issue carplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus extrhematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapputic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a human TERC-targeted double-stranded

Gaps

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26-FEB-2004

ADF93291;

RESULT 199

ઠે g

SiNA.

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explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human TERC-targeted double-stranded siNA, which is identical to the c-fos transcript target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; sinA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shrap, expression medulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human TERC transcript target sequence/siNA upper strand, SEQ ID 19.
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                                                                                                                                                                                                                                                                                                                                 Score 19; DB 1; Length 19;
Pred. No. 1.5e+02;
9; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                Sequence 19 BP; 3 A; 4 C; 3 G; 0 T; 9 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                           32 TGGCCATTTTTGTCTAAC 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 UGGCCAUUUUUUGUCUAAC 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP.
                                                                                                                                                                                                                                                                                                                               ch 4.2%;
1 Similarity 52.6%;
10; Conservative
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2002US-0386782P.
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2002US-0409293P.
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2002US-0406784P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADF93302 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
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09-SEP-2002; 2
15-JAN-2003; 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shrNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                        Gaps
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                                                                                       DB 1; Length 19;
1.5e+02;
                                                                                                                                      0; Indels
                                      Sequence 19 BP; 0 A; 3 C; 13 G; 0 T; 3 U; 0 Other;
                                                                                                      100.0%; Prec. ....
                                                                                       Score 19;
Pred. No.
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2002US-0363124P.
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                                                                                       4.28;
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                                                                           Query Match
Best Local Similarity 100.0
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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17-JUL-2002;
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05-SEP-2002;
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28-AUG-2003

X S X S X S X S X S S X S S X S S X S S X S S X S S X

WPI; 2003-689777/65.

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be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the sings include short interfering RNA (sirRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (sirRNA). The sinAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, and vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of sinA, conjugates and vector or shank, and vectors that express sin calls, tissue used to modulate expression of the telomerase genes in calls, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and used for treating cancer, restencesis, infectious diseases (specifically protozoal), transplant rejection, or autoimmume or age-related diseases, used for treating cancer, restencesis, infectious diseases (specifically skin ulcers and rheumatoid arthritis. The sinAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human TERC-targeted double-stranded sinA, which is identical to the c-fos transcript target sequence.
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100.0%; Pred. No. 1.5e+02;
tive 0; Mismatches 0; Indels
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2002US-0406784P.
2002US-0408378P.
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2002US-0386782P.
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Matches 19; Conservative
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05-SEP-2002;
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense of ligonuclectide. Specifically, the siNAs include short interfering RNA (shRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (shRNA). The siNAs can be unmodified or chemically synthesised, can contain deoxyribonuclectides, and can be chemically synthesised, can contain deoxyribonuclectides, and can be chemically synthesised, can contain deoxyribonuclectides, and can be chemically synthesised, can complaxes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue candor complaxes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue caplants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases. Ce.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The sinAs are also useful for drug careering, pharmacogenomics, strudying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence ceptors is indentical to the c-foe transcript target sequence.
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                                          New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                           Example 3; SEQ ID NO 7; 145pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 UGGGCCUGGGAGGGGUGGU 19
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nes 15; Conservative
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WO2003070742-A1.
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1D ADF93
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                                                                                                                                                                                                                                                                    The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise sense and antisense coligonaclectide. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA, and vectors that express sinA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer restenosis, infectious diseases (specifically, protozoal), transplant rejection, or autoimmune or age-related diseases, sin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic empaping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human TERC-targeted double-stranded siNA, which is identical to the c-fos transcript target sequence.
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                                                                                                                                                                                             New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                                                                                                                                                                Example 3; SEQ ID NO 12; 145pp; English.
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                 29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
2002US-0396600P.
                                                               15-JAN-2003; 2003US-0440129P
                                                                                                 (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                 Mcswiggen J, Beigelman L;
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Best Local Similarity 73.7
Matches 14; Conservative
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   17-JUL-2002;
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short contribering RNA (siRNA). double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically spothesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can complexes of siNA; and vectors that express in relates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue caplants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug correnning, diagnosis, lupus erythematosus, AIDS, macular degeneration, cannot in genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence consumption of a human TERC-targeted double-stranded
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                                                                                                                                                                                            11-MAR-2002; 2002US-0363124P.
06-UTV-2002; 2002US-0386782P.
17-UTL-2002; 2002US-0396600P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0408378P.
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                                                                            11-FEB-2003; 2003WO-US004088
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ses 19; Conservative
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28-AUG-2003
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Matches
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ADF93561 standard; RNA; 19 BP

26-FEB-2004 (first entry)

ADF93561;

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribourcleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an intisense oligonucleotide. Specifically, the siNAs include short hairpin RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, and/or complexes of siNA, and vectors that express siNA. The siNAs are and/or complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue caplants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for trasting cancer, restenosis, infectious diseases (specifically printiple sclerosis, lupus erythematosus, AlbS, macular degeneration, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AlbS, macular degeneration, cancer and validation, cancer in calls, therappate identification and validation, cancer in caper in calls.
Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; slNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 3; SEQ ID NO 285; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2002US-0396600P.
2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
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2002US-0363124P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mcswiggen J, Beigelman L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-689777/65.
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29-AUG-2002; 2
05-SEP-2002; 2
09-SEP-2002; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11-MAR-2002;
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The invention relates to short interfering nucleic acids (sinA) which downregulate expression of the one or more telomerase genes by RNA downregulate expression of the one or more telomerase genes by RNA be double or single stranded. They further comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the sinAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The sinAs can be unmodified or charmacontain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protected).
                                                                                                                                                            Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; double-stranded RNA; micro-RNA; miRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERT; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                                       Human TERC sina lower strand, SEQ ID 288.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2002US-0363124P.
2002US-03867B2P.
2002US-0396600P.
2002US-0406784P.
2002US-0408378P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2002US-0358580P
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06-JUN-2002;
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Sequence 19 BP; 3 A; 9 C; 7 G; 0 T; 0 U; 0 Other;

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Gaps

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Length 19;

4.2%; Score 19; DB 1; Length 19; 100.0%; Pred. No. 1.5e+02; ive 0; Mismatches 0; Indels

266 CCGGGGCTTCTCCGGAGGC 284

8

Conservative

Local Similarity nes 19; Conserv

Query Match Best Loca Matches 19 CCGGGGCTTCTCCGGAGGC 1

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonuclectide and an antisense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised. Expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue used to modulate expression of the telomerase genes in cells, tissue transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA, double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; siRNA, expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human TERC transcript target sequence/sinA upper strand, SEQ ID 14.
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                  Score 19; DB 1; Length 19;
Pred. No. 1.5e+02;
                                                                                   0; Indels
4.2%; bcc_
100.0%; Pred. No. ...
'... 0; Mismatches
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                                                                                                                                                            320 CCGCGGGTCTCTCGGGGC 338
                                                                                                                                                                                                        CCGCGGGTCTCTCGGGGGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-FEB-2002; 2002US-0358580P.
11-MAR-2002; 2002US-0363124P.
06-JUN-2002; 2002US-0386782P.
17-JUL-2002; 2002US-0396600P.
29-AUG-2002; 2002US-0466784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0408378P.
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                                                                                           Conservative
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                                                      Local Similarity
nes 19; Conserv
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                         Query Match
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protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, ALDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene rapping e.g., of single nucleoride polymorphisms). The present sequence represents the upper strand of a human TERC-targeted double-stranded siNA, which is identical to the c-fos transcript target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering uncleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short interfering RNA; shrap; and modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human TERC transcript target sequence/siNA upper strand, SEQ ID 26.
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                                                                                                                                                                                                          Score 19; DB 1; Length 19;
Pred. No. 1.5e+02;
5; Mismatches 0; Indels
                                                                                                                                                                           Sequence 19 BP; 2 A; 9 C; 3 G; 0 T; 5 U; 0 Other;
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                                                                                                                                                                                                                                                                                         140 GCCGCCTTCCACCGTTCAT 158
                                                                                                                                                                                                                                                                                                             1 GCCGCCUUCCACCGUUCAU 19
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2002US-0363124P.
2002US-0386702P.
2002US-0406784P.
2002US-0409233P.
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                                                                                                                                                                                                                              Local Similarity 73.7
Les 14; Conservative
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05-SEP-2002; 2
09-SEP-2002; 2
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06-JUN-2002;
17-JUL-2002;
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                                                                                                                                                                                                                                                                                                                                                                                      RESULT 207
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interfering RNA (BIRNA), double-stranded RNA, micro-RNA (miRNA) and short hairin RNA (BhRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNas are used to modulate expression of the telomerase genes in cells, tissue capants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restencies, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, ce.g. multiple sclerosis, lupus erythematosus AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation,
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                                                                                                                                                                                                                                                                                       genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human TERC-targeted double-stranded
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                                                                                                                                                                                                                                                                                                                                                    sinA, which is identical to the c-fos transcript target sequence
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2002US-0386782P.
2002US-0396600P.
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2002US-0408378P
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15-JAN-2003; 2003US-0440129P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mcswiggen J, Beigelman L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-689777/65.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO2003070742-A1.
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29-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-MAR-2002;
06-JUN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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New short interfering nucleic acid downregulates expression of the

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA convergulate expression of the one or more telomerase genes by RNA convergulate expression of the one or more telomerase genes by RNA convergence. The siNAs may or may not comprise sense and antisense coligonaclectide. They further comprise sense and antisense coligonaclectides, Specifically, the siNAs include and antisense oligonaclectides, Specifically, the siNAs include short interfering RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised. The invention also expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express sinA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue cused to modulate expression of the telomerase genes in cells, tissue cused to modulate expression of the telomerase genes in cells, tissue cused to modulate expression of the vivo gene therapy), or in grafts and transplants for every expressed in cells, tissue cused for treatenne or age-related diseases. Every in ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, so mapping (e.g., of single nuclectide polymorphisms). The present sequence crepresents the upper strand of a human TERC-targeted double-stranded cush, which is identical to the c-fos transcript target sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cytostatic, vasotropic, protozoacide, immunosuppressive, dermatological, neuroprotective, anti-HIV, ophthalmological; antiulcer, antirheumatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; mitco-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4.2%; Score 19; DB 1; Length 19; 94.7%; Pred. No. 1.5e+02; ve 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 19 BP; 5 A; 6 C; 7 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1; Mismatches
                                         Example 3; SEQ ID NO 27; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     374 AGGAACGGAGCGAGTCCCC 392
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 AGGAACGGAGCGAGUCCCC 19
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2002US-0386782P.
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2002US-0406784P.
2002US-0408378P.
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Matches 18; Conservative
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29-AUG-2002;
05-SEP-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 209
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA characterace. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense oligonacleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA). The siNAs can be unmodified or chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, a rautoimmune or age-related diseases, e.g. multiple sclerosis, lugue erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug correcting, diagnosis, therapeutic target identification and validation, screening, diagnosis, therapeutic target identification and validation, caption or sind, which is identical to the c-fost transcript target sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human TERC transcript target sequence/siNA upper strand, SEQ ID 15.
                                                                                                                                                                                                                       telomerase gene useful e.g. for treatment and diagnosis of cancer
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                                                                                                                                                                                                  New short interfering nucleic acid downregulates expression of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4.2%; Score 19; DB 1; Length 19; 84.2%; Pred. No. 1.5e+02; tive 3; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19 BP; 5 A; 5 C; 6 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                   Example 3; SEQ ID NO 9; 145pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 CCCUAACUGAGAAGGGCGU 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADF93298 standard; RNA; 19 BP
09-SEP-2002; 2002US-0409293P.
15-JAN-2003; 2003US-0440129P.
                                                                   (RIBO-) RIBOZYME PHARM INC.
                                                                                                             Mcswiggen J, Beigelman L;
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Best Local Similarity 84.2%
                                                                                                                                                         WPI; 2003-689777/65.
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the inversious traders to short interferring fluctures delicated by RNA communications. The single stranded. They further comprise sense and antisense interference. The single stranded. They further comprise sense and antisense condule or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotides. Specifically, the single include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short can contain deoxyribonucleotides, and can be chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised. Contains a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of single secondary complexes of single, by ex vivo gene therapy), or in grafts and cransplants for the treatment of a variety of conditions. They may be completed for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, conditions and rheumatoid arthritis. The sinks are also useful for drug screening, diagnosis, therapeutic target identification and validation, conspining (e.g., of single mucleotide polymorphisms). The present sequence conspining the upper strand of a human TERC-targeted double-stranded construction and validation, conspining, which is identical to the c-foe transcript target sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to short interfering nucleic acids (siNA) which
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4.2%; Score 19; DB 1; Length 19;
Best Local Similarity 84.2%; Pred. No. 1.5e+02;
Matches 16; Conservative 3; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 3; SEQ ID NO 15; 145pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 UUCUAGAGCAAACAAAAA 19
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                                                                                                                                                                                                                                                      17-JUL-2002; 2002US-0396600P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
                                                                       11-FEB-2003; 2003WO-US004088.
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2002US-0386782P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                   20-FEB-2002;
11-MAR-2002;
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28-AUG-2003
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ID ADF9
XX AC ADF9
XX ADF9
XX
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Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological;

Human TERC transcript target sequence/siNA upper strand, SEQ ID 30

26-FEB-2004 (first entry)

neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarbritic; antiaflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short interfering RNA; shraw, couple-stranded RNA; micro-RNA; miRNA; draw short hairpin RNA; shraw, tereasion modulation; gene therapy; dray screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense colligonacleotide. Specifically, the siNAs include short interfering RNA (shRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and example for treatment of a variety of conditions. They may be used for treatment of a variety of conditions. They may be used for treatment expenses; inpute erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, spentic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human TERC-targeted double-stranded crepted by which is identical to the c-fos transcript target sequence.
RNA interference; short interfering nucleic acid; siNA;
short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA;
short hairpin RNA; shRNA; expression modulation; gene therapy;
drug screening; diagnosis; therapeutic target identification;
pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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Pred. No. 1.5e+02;
2; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19 BP; 1 A; 9 C; 7 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 3; SEQ ID NO 28; 145pp; English.
                                                                                                                                                                                                                                                                                                                            2002US-0386782P.
2002US-0396600P.
2002US-0406784P.
2002US-0408378P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mcswiggen J, Beigelman L;
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Best Local Similarity
                                                                                                                                                                WO2003070742-A1.
                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                               11-FEB-2003;
                                                                                                                                                                                                                                                                                        20-FEB-2002;
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05-SEP-2002;
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06-JUN-2002;
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Gaps
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                392 CGCGCGCGCGCGATTCCC 410
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CGCGCGCGCGCGAUCCC 19

ò 셤 ADF93313 standard; RNA; 19 BP.

RESULT 212

ADF93313 ID ADF9

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonuclectide and an interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised. Can contain deoxyribonuclectides, and can be chemically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue cused to modulate expression of the telomerase genes in cells, tissue cused to modulate expression of the telomerase genes in cells, tissue camplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically crotozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug soreening, diagnosis, therapeutic target identification and sone mapping (e.g., of single nucleotide polymorphisms). The present sequence mapping (e.g., of single nucleotide polymorphisms). The present extranded contained the sequence contained the upper strand of a human TRRC-targeted double-etranded diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   siNA, which is identical to the c-fos transcript target sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 19; DB 1; 1
Pred. No. 1.5e+02;
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Best Local Similarity
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New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.

2002US-0363124P 2002US-0386782P 2002US-0406784P 2002US-0408738P 2002US-0408378P

17-JUL-2002; 29-AUG-2002; 2003US-0440129P

05-SEP-2002; L5-JAN-2003; (RIBO-) RIBOZYME PHARM INC

Mcswiggen J, Beigelman L; WPI; 2003-689777/65.

11-FEB-2003; 2003WO-US004088

20-FEB-2002;

11-MAR-2002; 06-JUN-2002;

WO2003070742-A1. Homo sapiens

28-AUG-2003

US09436060A-16.rng.sl

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue used to modulate expression of the telomerase genes in cells, tissue transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically cortozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, ekin ulcers and rheumatoid arthritis. The siNAs are also useful for drug
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  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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0; Indels
2; Mismatches
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                                                                             1 Acccaddacucdecucaca 19
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2002US-0406784P.

2002US-0408378P.

2002US-0409293P.

2003US-0440129P.
                                     428 ACCCAGGACTCGGCTCACA
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2002US-0363124P.
2002US-0386782P.
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                                                                                                                                                                                ADF93557 standard; RNA; 19
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                                                                                                                                                                                                                                                              (first entry)
  17; Conservative
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05-SEP-2002;
09-SEP-2002;
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17-JUL-2002;
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                                                                                                                                          RESULT 213
  Matches
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            genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a human TERC-targeted double-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 vasotropic; protozoacide; immunosuppressive; dermatological;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short, siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
screening, diagnosis, therapeutic target identification and validation,
                                                                                                                                                                      Gaps
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                                                                                                                                      Length 19;
                                                                                                                                                                      0; Indels
                                                                                                                                      DB 1; Le
1.5e+02;
                                                                                                  Sequence 19 BP; 2 A; 9 C; 7 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                        0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Human TERC siNA lower strand, SEQ ID 292.
                                                                                                                                      Score 19;
Pred. No.
                                                                                                                                                                                                      248 CTGGAGGCCGCGGTCGGCC 266
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2002US-0363124P.
2002US-0386782P.
2002US-0406784P.
2002US-0409378P.
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                                                                                                                                                                                                                                                                                                                                ADF93565 standard; RNA; 19
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                                                                                                                                      Query Match 4.2
Best Local Similarity 100.
Matches 19; Conservative
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05-SEP-2002;
09-SEP-2002;
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06-JUN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cytostatic;
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                                                                                                                                                                                                                                                                                              RESULT 214
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expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of sinA; conjugates and/or complexes of sinA, and vectors that express sinA. The sinAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restencesis, infections diseases (e.g. multiple sclerosis, invertice of conditions. They may be protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siMAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a human TERC-targeted double-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; pene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; couble-stranded RNA; micro-RNA; miRNA; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
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Pred. No. 1.5e+02;
                                                                                                                                                                                                                                                                                                                                                                         0; Indels
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100.0%; Pred. No. ...
... 0; Mismatches
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29-AUG-2002; 2002US-04067B4P-
0S-SEP-2002; 2002US-0408378P-
0S-SEP-2002; 2002US-0409293P-
1S-JAN-2003; 2003US-0440129P-
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2002US-0386782P.
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Best Local S
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Example 3; SEQ ID NO 278; 145pp; English.

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an interfering RNA (siNAA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (siNAA). Adouble-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised.

Expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be
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                                                                                                                                                                                                                                                                                                                                  used for treating cancer, restenosis, infectious diseases (apportitally protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, appendix engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a human TERC-targeted double-stranded
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100.0%; Pred. No. 1.5e+02;
ive 0; Mismatches 0;
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2002US-0363124P.
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2003US-0440129P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADF93293 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-FEB-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
les 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO2003070742-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-FEB-2002;
11-MAR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-JUN-2002;
17-JUL-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            39-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-JAN-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADF93293;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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CC an contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised.

CC an contain deoxyribonucleotides, and can be chemically synthesised.

CC an obtain RNA (siRNA). The siNAS can be unmodified or chemically modified, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are cused to modulate expression of the telomerase genes in cells, tissue cused to modulate expression of the telomerase genes in cells, tissue cused to readening (e.g., by ex vivo gene therapy), or in grafts and cranaplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, inputs erythematosus, finds are also useful for drug screening, diagnosis, therapeutic target identification and validation, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, and gene companing (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human TERCATEGET double-stranded company of the c-fos transcript target sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hiterfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hiterfering social; sindiagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                       New short interfering nucleic acid downregulates expression of the
                                                                                                                         relomerase gene useful e.g. for treatment and diagnosis of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4.2%; Score 19; DB 1; Length 19; 63.2%; Pred. No. 1.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19 BP; 1 A; 5 C; 6 G; 0 T; 7 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human TERC siNA lower strand, SEQ ID 273.
                                                                                                                                                        Example 3; SEQ ID NO 10; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               68 TAGGCGCCGTGCTTTTGCT 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 uAGGCGCCGUGCUUUGCU 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADF93546 standard; RNA; 19 BP
(RIBO-) RIBOZYME PHARM INC.
                                  Mcswiggen J, Beigelman L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-FEB-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity 63.2
nes 12; Conservative
                                                                    WPI; 2003-689777/65.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADF93546/
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The inversion between the one or more telomerase genes by Nun. Computed and control stands (stranded They further comprise sense and antisense regions, or alternatively are assembled from a sense onlique stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siNAA), double-stranded RNA, micro-RNA (miRNA) and short can contain deoxyribonucleotides, and can be chemically synthesised, can complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue cand/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue carpaisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be concerned to the treatment of a variety of conditions. They may be concerned to the treatment of a variety of conditions. They may be concerned to the treatment of a variety of conditions. They may be concerned to the treatment of a variety of conditions. They may be concerned to the treatment of a variety of conditions of secreening dependence of single mucleotide polymorphisms). The present sequence can mapping (e.g., of single mucleotide polymorphisms). The present sequence concerned the lower strand of a human TERC-targeted double-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering conclet acid; siNA; short hairpin RNA; shRNA; expression modulation; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to short interfering nucleic acids (siNA) which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 19 BP; 3 A; 6 C; 5 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Pred. No. 1.5e+02; ive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 3; SEQ ID NO 273; 145pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 50 CCCTAACTGAGAGGGCGT 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19 CCCTAACTGAGAAGGGCGT 1
                                 2002US-0363124P.
2002US-0386782P.
2002US-0396600P.
2002US-0406784P.
2002US-0408378P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADF93562 standard; RNA; 19 BP
                                                                                                                                                                                                                    09-SEP-2002; 2002US-0409293P.
15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                                                                                                                             (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                    Mcswiggen J, Beigelman L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-689777/65.
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20-FEB-2002;
11-MAR-2002;
06-JUN-2002;
17-JUL-2002;
29-AUG-2002;
05-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26-FEB-2004
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ADF93562/c
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11-FEB-2003; 2003WO-US004088.

(first entry)

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Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HTV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shrNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                          Human TERC transcript target sequence/sinA upper strand, SEQ ID 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                    11-FEB-2003; 2003WO-US004088
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mcswiggen J, Beigelman L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-689777/65.
                                                                                                                                                                                                                                                                                                                                            WO2003070742-A1.
                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-JUN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-AUG-2002;
05-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-FEB-2002;
26-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-MAR-2002;
                                                                                                                                                                                                                                                                                                                                                                                         28-AUG-2003
The invention relates to short interfering nucleic acids (siNA) which downtregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in calls, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and trensplants for the treatment of a variety of conditions. They may be used for treating cancer, restenoisis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lubus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthities. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, agenetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence
                          88
drug screening, diagnosis, therapeutic target identification;
pharmacogenomics, gene function analysis, gene mapping; TERC; TERT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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Pred. No. 1.5e+02;
0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19 BP; 2 A; 9 C; 5 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 3; SEQ ID NO 289; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4.2%; Scc...
100.0%; Prev
                                                                                                                                                                                                                                                                11-MAR-2002; 2002US-0363124P.
06-JUN-2002; 2002US-0386782P.
17-JUL-2002; 2002US-0396600P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0408378P.
                                                                                                                                                                                                     11-FEB-2003; 2003WO-US004088
                                                                                                                                                                                                                                                                                                                                                                                                                                                            (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mcswiggen J, Beigelman L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 19; Conservat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-689777/65.
                                                                                                             402003070742-A1
                                                                    Homo sapiens.
                                                                                                                                                          28-AUG-2003
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2002US-0386782P. 2002US-0396600P. 2002US-0406784P. 2002US-0408378P.

2002US-0363124P

2002US-0409293P 2003US-0440129P

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, can condulate expression of the telomerase genes in cells, tissue and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue candot the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically proteoxoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AlbS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, expresents the upper strand of a human TERC-targeted dubble extremeded mapping (e.g., of single nucleotide polymorphisms). The present extranded mapping (e.g., of single nucleotide polymorphisms). The present sequence extranded dependence and rheumatoid a human TERC-targeted dubble extranded degeneration and sense extranded deserved and the upper strand of a human TERC-targeted dubble extranded degeneration and sense extranged degeneration and sense extranged degeneration and sense extranged and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4.2%; Score 19; DB 1; Length 19;
89.5%; Pred. No. 1.5e+02;
tive 2; Mismatches 0; Indels
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Example 3; SEQ ID NO 25; 145pp; English.
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Best Local Similarity 89.54
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Gaps

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338 CGAGGCCGAGGTTCAGGCC 356

Conservative

CGAGGGGAGGTTCAGGCC 1

13

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ADF93308 standard; RNA; 19 BP

RESULT 219

ADF93308

ADF93308 ID ADF9 XX AC ADF9 XX

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Gape

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338 CGAGGGGAGGTTCAGGCC 356

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Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
                                                                                                                            Human TERC siNA lower strand, SEQ ID 279.
BP.
                                                                                                                                                                                                                                                                                                                                                                 17-JUL-2002; 2002US-0396600P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
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2002US-0386782P.
                                                                                                                                                                                                                                                                                                                                                                                                09-SEP-2002; 2002US-0409293P.
                                                              ADF93552 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                 (RIBO-) RIBOZYME PHARM INC.
                                                                                                         (first entry)
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                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                         26-FEB-2004
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06-JUN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mcswiggen J,
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                                                                                    ADF93552;
                                           RESULT 220
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonuclectide and antisense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates confaint of stressed from the tenderses genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and candor complants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplants rejection, or autoimmune or age-related diseases of sind, diagnosis, therapeutic carget identification and validation, can under emphing (e.g., of single nucleoride polymorphisms). The present sequence mapping (e.g., of single nucleoride polymorphisms). The present sequence Example 3; SEQ ID NO 279; 145pp; English.

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                                                                                                                                                                                                                                                                                                                                                                                                             Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerses; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering nucleic acid; siNA; short interfering RNA; alkNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shrNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                       Gaps
represents the lower strand of a human TERC-targeted double-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New short interfering nucleic acid downregulates expression of the
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                                                                                       Length 19;
                                                                                                                      0; Indels
                                                 Sequence 19 BP; 3 A; 2 C; 3 G; 0 T; 11 U; 0 Other;
                                                                                     DB 1; Le
1.5e+02;
                                                                              Human TERC siNA lower strand, SEQ ID 293.
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                                                                                                                                                      158 TTCTAGAGCAAACAAAAA 176
                                                                                                                                                                          ADF93566 standard; RNA; 19 BP.
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2002US-0363124P.
2002US-0386782P.
2002US-0396600P.
2002US-0406784P.
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                                                                                                                                                                                                                                                                                                                                              26-FEB-2004 (first entry)
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                                                                                 Query Match
Best Local Similarity 100.
Matches 19; Conservative
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06-JUN-2002;
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05-SEP-2002;
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used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restencials, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence regressents the lower strand of a human TERC-targeted double-stranded
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human TERC transcript target sequence/siNA upper strand, SEQ ID 13.
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                                                                                                                                                                                                                                                         Sequence 19 BP; 4 A; 8 C; 4 G; 0 T; 3 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                            410 CTGAGCTGTGGGACGTGCA 428
                                                                                                                                                                                                                                                                                                                                                                                                    19 CTGAGCTGTGGGACGTGCA 1
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05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
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2002US-0386782P.
2002US-0396600P.
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06-JUN-2002;
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interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense cregions, or alternatively are assembled from a sense oligonucleotide specifically, the siNAs include short an antisense oligonucleotide. Specifically, the siNAs include short history (ShRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised. Commodiate expressed from a vector or enzymatically synthesised. The invention also relates to kits for the invitro or in vivo delivery of siNA, conjugates and/or complexes of siNA, and vectors that express siNA. The siNas are used to modulate expression of the telomerase genes in calls, tissue complexes of siNA, and vectors that express siNA. The siNas are used to modulate expression of the telomerase genes in calls, tissue complexes of siNA, and vectors that express siNA. The siNas are used for treating cancer, restencesis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, protozoal), transplant rejection, or autoimmune or age-related diseases, sin ultiple sclerosiss, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNas are also useful for drug screening, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human TERC-targeted double-stranded signal.
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89.5%; Pred. No. 1.5e+02;
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2; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        122 GCGGAAAAGCCTCGGCCTG 140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 GCGGAAAAGCCUCGGCCUG 19
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11-MAX-2002; 2002US-03631Z4P.
06-UJN-2002; 2002US-0386782P.
17-UUL-2002; 2002US-0386600P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0409378P.
09-SEP-2002; 2002US-0409233P.
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Best Local Similarity 89.5
Matches 17; Conservative
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Mcswiggen J, Beigelman L;

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and matisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA, and vectors that express since it since are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restencisis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus. AIDS, macular degeneration, skin ulcers and rheumatosid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, agenematic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a human TERC-targeted double-stranded
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                                                              New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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100.0%; Pred. No. 1.5e+02;
ive 0; Mismatches 0; Indels
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                                                                                                                                    Example 3; SEQ ID NO 271; 145pp; English.
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11-MAR-2002; 2002US-0363124P.
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                WPI; 2003-689777/65
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Best Local Similarity
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (shRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain decoxribbonucleotides, and can be themically synthesised, can contain decoxribbonucleotides, and can be themically synthesised, can contain decoxribbonucleotides, and can be themically synthesised, can contain decoxribbonucleotides, and vectors that express sinA. The siNAs are cused to modulate expression of the telomerase genes in cells, tissue cand/or complexes of siNA, and vectors that express siNA. The siNAs are cused to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be cused for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, ce.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening diagnosis, therapeutic target identification and validation, mapping (e.g., of single mucleotide polymorphisms). The present sequence mapping (e.g., of single mucleotide polymorphisms). The present sequence contains the lower strand of a human TERC-targeted double-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthitic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNN; micro-RNA; miRNA; short halpin RNA; shrNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                     New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4.2%; Score 19; DB 1; Length 19;
100.0%; Pred. No. 1.5e+02;
Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19 BP; 3 A; 5 C; 10 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human TERC siNA lower strand, SEQ ID 286.
                                                                                                                                                                                                                                                                                                                                                                                                   Example 3; SEQ ID NO 281; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          194 TTCGCCCCCCCGGGGACC 212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19 Tresecerceseseace 1
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                       17-JUL-2002; 2002US-0396600P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0409378P.
09-SEP-2002; 2002US-0409293P.
2002US-0386782P.
                                                                                                                          15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                           (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADF93559 standard; RNA; 19
                                                                                                                                                                                                                          Mcswiggen J, Beigelman L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity 100.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-FEB-2004 (first entry)
                                                                                                                                                                                                                                                                           WPI; 2003-689777/65.
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonuclectide and antisense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AlbS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, agenetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the lower strand of a human TERC-targeted double-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19 BP; 1 A; 3 C; 10 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 3; SEQ ID NO 286; 145pp; English.
                                                                                                                                                                                                       2002US-0396782P.
2002US-0396600P.
2002US-0406784P.
                                                                                                                                                                                                                                                                    2002US-0408378P.
                                                                                                                         11-FEB-2003; 2003WO-US004088
                                                                                                                                                                                    2002US-0363124P
                                                                                                                                                                                                                                                                                                              2003US-0440129P
                                                                                                                                                                                                                                                                                                                                                       (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                Mcswiggen J, Beigelman L;
                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-689777/65.
                                       WO2003070742-A1.
                                                                                                                                                                                                                                                                    05-SEP-2002; 2
09-SEP-2002; 2
15-JAN-2003; 2
  Homo sapiens.
                                                                                                                                                                                                            06-JUN-2002;
                                                                                                                                                                                                                           17-JUL-2002;
                                                                                                                                                                                                                                                   29-AUG-2002;
                                                                                                                                                                20-FEB-2002;
                                                                                                                                                                                  11-MAR-2002;
                                                                                28-AUG-2003
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Claim 12; SEQ ID NO 8; 58pp; English.

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                      Gaps
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Score 19; DB 1; Length 19;
Pred. No. 1.5e+02;
                    0; Indels
                                                                                                                                                                                 Human telomerase RNA gene, SDSO target region #4.
    100.0%; Pred. .v.
                                         284 CACCCACTGCCACCGCGAA 302
                                                              19 cacccacreccaccecaa 1
                                                                                                                  ADO22919 standard; cDNA; 19 BP
 4.28;
                                                                                                                                                            (first entry)
                     19; Conservative
 Query Match
Best Local Similarity
                                                                                                                                                            01-JUL-2004
                                                                                                                                       ADO22919;
                                                                                               RESULT 226
                     Matches
                                                                                                          AD022919
                                                              용
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Human; ss; SDSO; short double stranded oligonucleotide; cleavage site; viral infection; malignant tumour; genetic disease; metabolic disease; gene chip; protein chip; microarray; gene drug; Dermogene; Lungene; Hepatogene; Leukogene; Lymbhogene; Prostogene; Breastogene; Braintumogene; Skin-whitogene; short interfering RNA; siRNA; cancer;
                                                                                                                                                                                               Designing and selecting short double-stranded oligonucleotides for treating viral infections, cancer and genetic or metabolic diseases, comprises using gene chip and protein chip microarrays to identify
                                                                                                               16-SEP-2002; 2002US-00016490.
                                                                                                                               16-SEP-2002; 2002US-00016490.
                                                                                                                                                                                                                       specific DNA sequences.
                                                                                                                                                                               WPI; 2004-355427/33
                                              RNA interference
                                                                               US2004072769-A1
                                                                                                                                               (YINJ/) YIN J Q.
                                                               Homo sapiens
                                                                                               L5-APR-2004
                                                                                                                                                                Yin JQ;
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The invention relates to screening, identifying or predicting, and cassembling 19-25 in debuble-stranded oligomucleotides (termed short debuble stranded oligomucleotides) as active pharmaceutical compositions cassembling 19-25 in debuble stranded oligomucleotides (termed short debuble for the treatment of viral infections, malignant tumours, and genetic and metabolic diseases, comprising screening and identifying a specific DNA sequence in an abnormal gene encoding a protein with gene chip and protein chip microarrays, identifying a specific DNA sequence within the abnormal gene encoding a protein or playing other biological robe with the absistance of computer and specific software. Computering forces within the abnormal gene encoding a protein or playing other biological robe with the absistance of computer and specific software. Computering fitting from the standard specific software, computering the specific software. Computering fitting from the standard specific software. Al(7) CCG-3' or 5'-U(7) CCG-3' specific and making sure that selected sequence is not localised within the stem-loop of target mRNA with any related software. Also included are pharmaceutical compositions of gene drugs from of an RNA molecule and RNA, 19-25 in taken with the special pattern, 5'-CGGAT(0)-3' or its derivatives, one or more more nucleic acid condensation genes for none, one or more pharmaceutical carites, one or more specific cell-cargeting proteins and other active agents and additional materials) and a simplified method for predicting and selecting of specific and efficacious small double-stranded oligomucleotide molecules or short interfering she specifically of specific and a simplified method for predicting and selecting and selecting specific and activatives, one or short interfering and selecting and selecting secion of an oligomucleotide molecules or short interfering and selecting secion of an oligomucleotide sequence evaluating the specifically inactivate a group of corresponding genes in objections, cancer, or sp

Sequence 19 BP; 6 A; 4 C; 8 G; 1 T; 0 U; 0 Other;

4.2%; Score 19; DB 1; Length 19;

Query Match

Gaps ö Indels No. 1.5e+02; natches 0; 0; Mismatches 372 AGAGGAACGGAGCGAGTCC 390 100.0%; Best Local Similarity 100. Matches 19; Conservative 8 g

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1 AGAGGAACGGAGCGAGTCC 19

RESULT 227

ADO23063 standard; cDNA; 19 BP.

ADO23063;

(first entry) 01-JUL-2004 Human telomerase RNA gene, SDSO target region #3.

Human; ss; SDSO; short double stranded oligonucleotide; cleavage site; viral infection; malignant tumour; genetic disease; metabolic disease; gene chip; protein chip; microarray; gene drug; Dernogene; Lungene; Hepatogene; Lungene; Lungene; Lungene; Lungene; Lungene; Lungene; Lungene; Asin-whitogene; short interfering RNA; siRNA; cancer; RNA interference

Homo

US2004072769-A1.

15-APR-2004.

16-SEP-2002; 2002US-00016490.

16-SEP-2002; 2002US-00016490.

Q P NIX (/CNIX)

Yin JQ;

WPI; 2004-355427/33.

Designing and selecting short double-stranded oligonucleotides for treating viral infections, cancer and genetic or metabolic diseases, comprises using gene chip and protein chip microarrays to identify specific DNA sequences.

Example 1; Page 19; 58pp; English.

The invention relates to screening, identifying or predicting, and assembling 19-25 nt double-stranded oligonucleotides (termed short double stranded oligonucleotides, SDSO) as active pharmaceutical compositions for the treatment of viral infections, malignant tumours, and genetic and metabolic diseases, comprising screening and identifying a specific DNA sequence in an abnormal gene encoding a protein with gene chip and protein chip microarrays. The above method comprises screening the disease-causing genes, over-expressing in cells and/or tissues, with the abnormal gene encoding a protein or playing other comprised within the abnormal gene encoding a protein or playing other compositions of the protein or predicting efficacious 19-25 nt double-stranded oligonucleotides with a conformation of an RNA molecule and making sure that selected sequence is not localised within the stem-loop of target mRNA with any related software. Also included are pharmaceutical compositions of gene drugs (such as Dermogene, Dungone, Hepatogene, Loukogene, Lymphogene, Prostogene, Breastogene, Braintumogene and Skin-whitogene including but corous of specific 19-25 nt daRNA, 19-25 nt daRNA, and/or single-stranded RNA and/or NNA with the special pattern, 5'-cogaroup of specific 19-25 nt daRNA, 19-25 nt daRNA, and/or single-stranded RNA and/or NNA with the special pattern, 5'-cogaroup of specific la-25 nt daRNA, 19-25 nt daRNA, and/or single-stranded RNA and/or NNA with the special pattern, 5'-cogaroup of specific cell-targeting proteins and other active agents and additional materials) and a simplified method for predicting and selecting a meterials) and simplified method for predicting and selecting and additional

antisense oligonuclectide molecules or short interfering RNA (siRNA) (comprising identifying a special pattern that can be localised in any position of an oligonuclectide sequence evaluating the specificity of a selected sequence). The Short interfering RNA (siRNA) are targeted against genes involved in viral infection, malignant tumours, genetic and methodic diseases. The methods are useful for designing and selecting short double-stranded oligonucleotides as a gene drug that can specifically inactivate a group of corresponding genes. The composition may be used for treating diseases or disorders associated with abnormal expression of genes in cells or tissues of humans or animals, such as viral infections, cancer, or genetic or metabolic diseases. The present sequence is a target region for an SDSO from an human cDNA. specific and efficacious small double-stranded oligonucleotides (SDSO), \$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$\$

Sequence 19 BP; 3 A; 7 C; 6 G; 3 T; 0 U; 0 Other;

Gaps ., Score 19; DB 1; Length 19; Pred. No. 1.5e+02; 0; Indels Query Match 4.2%; Score 19; DB Best Local Similarity 100.0%; Pred. No. 1.5 Matches 19; Conservative 0; Mismatches

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RESULT 228

ADO23064 standard; cDNA; 19 BP

ADO23064;

01-JUL-2004 (first entry)

Human telomerase RNA gene, SDSO target region #5.

Human; ss; SDSO; short double stranded oligonucleotide; cleavage site; viral infection; malignant tumour; genetic disease; metabolic disease; gene chip; protein chip; mcroarray; gene drug; Dermogene; Lungene; Hepatogene; Leukogene; Lymbhogene; Prostogene; Breatogene; Braintumogene; Skin-whitogene; short interfering RNA; siRNA; cancer; RNA interference

Homo sapiens.

US2004072769-A1.

15-APR-2004

16-SEP-2002; 2002US-00016490.

16-SEP-2002; 2002US-00016490.

(YINJ/) XIN J Q.

Yin JQ;

WPI; 2004-355427/33.

Designing and selecting short double-stranded oligonucleotides for treating viral infections, cancer and genetic or metabolic diseases, comprises using gene chip and protein chip microarrays to identify specific DNA sequences.

Example 1; Page 19; 58pp; English.

The invention relates to screening, identifying or predicting, and assembling 19-25 nt double-stranded oligonucleotides (termed short double stranded oligonucleotides, SDSO) as active pharmaceutical compositions for the treatment of viral infections, malignant tumours, and genetic and metabolic diseases, comprising screening and identifying a specific DNA sequence in an abnormal gene encoding a protein with gene chip and protein chip microarrays. The above method comprises screening the disease-causing genes, over-expressing in cells and/or tissues, with the

WPI; 2004-355427/33.

agents (or none), one or more pharmaceutical carriers parcell, 3 conference and condensation agents (or none), one or more pharmaceutical carriers, one or more perity of perity of the second additional materials) and a simplified method for predicting and selecting and additional materials) and esimplified method for predicting and selecting and efficacious small double-stranded oligonucleotides (SDSO), antisense oligonucleotide molecules or short interfering RNA (siRNA) (comprising identifying a special pattern that can be localised in any possition of an oligonucleotide sequence evaluating the specificity of a selected sequence). The Short interfering RNA (siRNA) are targeted against genes involved in viral infection, malignant tumours, genetic and metabolic diseases. The methods are useful for designing and selecting short double-stranded oligonucleotides as a gene drug that can specifically inactivate a group of corresponding genes. The composition may be used for treating diseases or disorders associated with abnormal expression of genes in cells or tissues of humans or animals, such as viral infections, cancer, or genetic or metabolic diseases. The present sequence is a target region for an SDSO from an human cDNA. gene chip and protein chip microarrays, identifying a specific DNA sequence within the abnormal gene encoding a protein or playing other biological roles with the assistance of computer and specific software, predicting efficacious 19-25 int double-stranded oligonucleotides with a 5'-AU(T)CCG-3' or 5'-U(T)CCCG-3' special pattern complementary to at least a portion of an RNA molecule and making sure that selected sequence is not localised within the stem-loop of target mRNA with any related software. Also included are pharmaceutical compositions of gene drugs (such as Dermogene, Lungene, Hepatogene, Leukogene, Lymphogene, Braintunogene and Skin-whitogene including but being not limited to part or all of the following components: single or a group of specific 19-25 it daRNA, 19-25 it sRNA-CDNA, 19-25 it daRNA, or single or a single or s

Sequence 19 BP; 1 A; 2 C; 12 G; 4 T; 0 U; 0 Other;

Score 19; DB 1; Length 19; Pred. No. 1.5e+02; 0; Mismatches 0; Indels 4.2. 100.0%; Pr. 14 TGGGCCTGGGAGGGGTGGT 32 1 receccrecasederec 19 Query Match
Best Local Similarity 100.
Warches 19; Conservative 용 ò

Gaps ..

> ADO23062 standard; cDNA; 19 BP ADO23062; RESULT 229 AD023062 XXX AC AD02 XXX XXX XXX XXX XXX HUMB KW HOPS KW Brai KW Brai KW Brai KW Brai KW Brai KW Brai KW HOPS KW

Human telomerase RNA gene, SDSO target region #2. 01-JUL-2004 (first entry)

Human, 88; SDSO; short double stranded oligonucleotide; cleavage site; viral infection; malignant tumour; genetic disease; metabolic disease; gene chip; protein chip; microarray; gene drug; Dermogene; Lungene; Hepatogene; Lunkogene; Lymphogene; Prostogene; Brastogene; Braintumogene; Skin-whitogene; short interfering RNA; siRNA; cancer; RNA interference

Homo sapiens.

US2004072769-A1.

15-APR-2004.

16-SEP-2002; 2002US-00016490.

16-SEP-2002; 2002US-00016490.

(YINJ/) YIN J Q.

Yin JQ;

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The invention relates to screening, identifying or predicting, and casemaling 19-25 in double-stranded oligomucleotides (termed short double stranded oligomucleotides) as acrive pharmaceutical compositions for the treatment of viral infections, malignant tumours, and genetic and for the treatment of viral infections, malignant tumours, and genetic and control of oligomucleotides with a sequence in an abnormal gene encoding a protein with gene chip and protein chip microarrays. The above method comprises screening the protein chip microarrays. The above method comprises screening the protein chip microarrays. The above method comprises screening the protein chip microarrays. The above method comprises screening the protein chip microarrays in cells and/or tissues, with the case and protein chip microarrays, identifying a specific DNA sequence within the abnormal gene encoding a protein or protein chip with the assistance of computer and specific DNA candidated within the aboremal gene encoding a protein or protein of an RNA molecule and making sure that selected sequence of predicting efficacious 19-25 nt double-stranded oligomucleotides with a 5'-AU(T)CCCG-3' or 5'-U(T)CCCG-3' special pattern complementary to at least a portion of an RNA molecule and making sure that selected sequence of such or localised within the stem-loop of target make with a sector of such or localised within the stem-loop of target make with a prediction of such as Dermogene, inagene, Hapatogene, Leukogene, Induke or molecule of sector of such as Dermogene, Dara and/or DNA with the special pattern, 5'-CCGMT(U)-3' or its derivatives, one or more pacific cell-targeting proteins and other active agents and additional capents of an encourage of section of an oligomucleotide molecule sequence evaluating the specific and efficacious small double-erranded oligomucleotide sequence evaluating the specific and efficacious small double-erranded oligomucleotide sequence evaluating the specific and efficacious small development of against genes invol
                                                 Designing and selecting short double-stranded oligonucleotides for treating viral infections, cancer and genetic or metabolic diseases, comprises using gene chip and protein chip microarrays to identify specific DNA sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequence is a target region for an SDSO from an human cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19 BP; 5 A; 6 C; 7 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                 Example 1; Page 19; 58pp; English.
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DB 1; Length 19; 1.5e+02; 0; Indela A.2%; Score 19; DB Local Similarity 100.0%; Pred. No. 1.5 les 19; Conservative 0; Mismatches 116 CAGCGGGGGGAAAAGCCTC 134 1 cagcegeceananacere 19 Query Match Matches ઠે 셤

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Gaps

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Human telomerase RNA gene, SDSO target region #6. (first entry) 01-JUL-2004 AD023065;

ADO23065 standard; cDNA; 19

RESULT 230

AD023065

Human; 88; SDSO; short double stranded oligonucleotide; cleavage site; viral infection; malignant tumour; genetic disease; metabolic disease; gene chip; protein chip; microarray; gene drug; Dermogene; Lungene; Hepatogene; Leukogene; Lymphogene; Prostogene; Breastogene;

1 CCAGCCCCCGAACCCCGCC 19

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Braintumogene; Skin-whitogene; short interfering RNA; siRNA; cancer;
                                        WPI; 2004-355427/33.
  RNA interference
            US2004072769-A1.
                               (YINJ/) YIN J Q.
       Homo sapiens.
                 15-APR-2004.
                                    7in JQ;
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26-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                03-JUN-2004.
                                                                                                ADP27906;
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Matches
RESULT 2
ADP27906
                                                                                                The invention relates to screening, identifying or predicting, and sembling 19-25 nt double-stranded oligonucleotides (termed short double stranded oligonucleotides, SDSO) as active pharmaceutical compositions cranded oligonucleotides, SDSO) as active pharmaceutical compositions cranded oligonucleotides, SDSO) as active pharmaceutical compositions for the treatment of viral infections, malignant tumours, and genetic and metabolic diseases, comprising screening and identifying a specific DNA sequence in an abnormal gene encoding a protein with gene chip and protein chip microarrays. The above method comprises screening the disease-causing genes, over-expressing in cells and/or tissues, with the disease-causing genes, over-expressing, identifying a specific DNA sequence within the assistance of computer and specific DNA composition of an RNA molter encoding a protein or playing other biological roles with the assistance of computer and specific DNA composition of an RNA molter encoding a protein or playing other composition of an RNA molter encoding a protein or playing other composition of an RNA molter encoding a protein or playing other composition of an RNA molter encoding a protein of gene drugs of the seatogene, branched are pharmaceutical compositions of gene drugs of specific 19-25 in dsRNA, 19-25 in dsRNA, 19-25 in dsRNA and or single or more pharmaceutical compositions of gene (or none), one or more pharmaceutical carriers, one or more pharmaceutical carriers of comparing deficient of a simplified method for praceficing and encoded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         expression of genes in cells or tissues of humans or animals, such as viral infections, cancer, or genetic or metabolic diseases. The present
                                                                                                                                                                                                                                                                                                                                                                                      Designing and selecting short double-stranded oligonucleotides for treating viral infections, cancer and genetic or metabolic diseases, comprises using gene chip and protein chip microarrays to identify
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        viral infections, cancer, or genetic or metabolic diseases. sequence is a target region for an SDSO from an human CDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 19; DB 1; Length 19;
Pred. No. 1.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19 BP; 3 A; 13 C; 3 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4.2%; Scor.
100.0%; Pred. No. 1...
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                                                                                                                                                                                                 16-SEP-2002; 2002US-00016490.
                                                                                                                                                   16-SEP-2002; 2002US-00016490.
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Best Local Similarity 100.
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           specific DNA sequences.
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This invention relates to a novel method concerning prognostic markers associated with EGFR (epidermal growth factor receptor) positive cancer. Specifically, it refers to a gene expression profiling method that can provide a prediction as to whether a patient is likely to respond well to treatment with an EGFR inhibitor. The present invention describes the currentiative analysis of the expression level of the RNA transcript of at CRP17, LAMC2 or their products thereof. It further provides a cDNA microarray containing named genes that represent prognostic transcripts which are useful for determining whether a patient diagnosed with an EGFR capressing head or neck cancer or colon cancer exhibits elevated or decreased expression levels of these genes compared to normal. As such, these methods are also useful for prognosing or predicting the likelihood or cancer-attributable death or prognession, including recurrence and metastatic spread of a neoplastic disease, as well as drug resistance.

This oligonucleotide sequence is a PCR primer used to amplify a human PCR amplify a such; and a prognostic cancer marker, given in an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Prognosing a patient with EGFR-expressing colon cancer comprises subjecting a sample comprising EGFR-expressing cancer cells to quantitative analysis of the expression level of the RNA transcript of at
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                                                                                                                                                                                                                                      PCR primer to amplify a human cancer prognostic marker DNA SeqID 343.
                                                                                                                                                                                                                                                                                                          human; primer; PCR; prognostic marker; BGPR; epidermal growth factor receptor; cancer; gene expression profiling; microarray; head and neck cancer; colon cancer; metastatic spread;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Baker JB, Cronin MT, Shak S, Baselga J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 54; SEQ ID NO 343; 113pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (VALL-) VALL HEBRON UNIV HOSPITAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     371 AAGAGGAACGGAGCGAGTC 389
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 AAGAGGAACGGAGCGAGTC 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               east one gene e.g., CD44v3.
ADP27906 standard; DNA; 19
                                                                                                                                                             (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                              neoplastic disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-420643/39.
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es 19; Conserv
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Gaps

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0; Indels

230 CCAGCCCCCGAACCCCGCC 248

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'note = "2',4'-oxyethylene linkage in the sugar residues"

OTHER

Location/Qualifiers

(first entry)

..5 *tag= a mod base= /mod_base= OTHER /note= "2',4'-oxyethylene linkage in the sugar residues"

/*tag= b /mod_base= OTHER /note= "C-hydroxyethyl phosphate"

Morita K;

/mod_base= OTHER /note= "phosphorothioate backbone' 15. 19

/*tag= c

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*tag=

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Stable 2',5'-oligoadenylic acid analogs containing natural and modified nucleic acid units as well as unusual phosphate groups with excellent activity particularly antitumor, applicable in cancer or antiviral
                                                                                                                                             Cytostatic; virucide; 2'; 5'-oligoadenylic acid analog; antitumour;
                                                                                     2',5'-oligoadenylic acid analog related oligonucleotide #4
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                                                                                                                                                                           antiviral; cancer; ss
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                                                                                                                                                                                                                          Synthetic.
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             This invention relates to a novel method concerning prognostic markers associated with EGFR (epidermal growth factor receptor) positive cancer. Specifically, it refers to a gene expression profiling method that can provide a prediction as to whether a patient is likely to respond well to treatment with an EGFR inhibitor. The present invention describes the quantitative analysis of the expression level of the RNA transcript of least one gene selected from the group of CD44v6, DBS, GRO1, KRT17, LAMC2 or their products thereof. It further provides a cDNA microarray containing named genes that represent prognostic transcripts which are useful for determining whether a patient diagnosed with an EGFR expressing head or neck cancer or colon cancer exhibits elevated or decreased expression levels of these genes compared to normal. As such, these methods are also useful for prognession, including recurrence and metastatic spread of a neoplastic disease, as well as drug resistance. This oligonucleotide sequence is a PCR primer used to amplify a human PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Prognosing a patient with EGFR-expressing colon cancer comprises subjecting a sample comprising EGFR-expressing cancer cells to quantitative analysis of the expression level of the RNA transcript of at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   amplicon DNA sequence used as a prognostic cancer marker, given in an exemplification of the invention.
                                                                                                                                                                                                 PCR primer to amplify a human cancer prognostic marker DNA SeqID 344.
                                                                                                                                                                                                                                              human; primer; PCR; prognostic marker; BGFR; expression profiling; epidermal growth factor receptor; cancer; gene expression profiling; microarray; head and neck cancer; colon cancer; metastatic spread;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4.2%; Score 19; DB 1; Length 19; 100.0%; Pred. No. 1.5e+02; Live 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Shak S, Baselga J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             54; SEQ ID NO 344; 113pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (GENO-) GENOMIC HEALTH INC.
(VALL-) VALL HEBRON UNIV HOSPITAL.
ADP27907/C

ADP27907 standard; DNA; 19 BP. XX
AC ADP27907;
XX
DI 26-AUG-2004 (first entry)
XX
DI 26-AUG-2004 (first entry)
XX
Whuman; primer; PCR; prognostic mana decreation of the pridermal growth factor receptor with microarray; head and neck cancer with microarray; head and neck cancer XX
XX
Homo sapiens.
XX
Homo sapiens.
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HOMO-2003; 2002US-0427090P.
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MO2004046386-A1.
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MO2004046386-A1.
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HOMO-2003; 2002US-0427090P.
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MO2004046386-A1.
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HOMO-2003; 2002US-0427090P.
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MPL; 2004-420643/39.
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GENO-) GENOMIC HEALTH INC.
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HOMO-2003; 2002US-0427090P.
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HOMO-2003; 2002US-0427090P.
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GENO-) GENOMIC HEALTH INC.
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HOMO-2003; 2003WO-US036777.
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GENO-) GENOMIC HEARTH OF PREADLY 
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                               The present invention relates to novel 2',5'-oligoadenylic acid analogs and their pharmacologically- acceptable salts. The analogs are stable with superior antitumour and antiviral activity and so are useful in cancer or antiviral therapy e.g. as antisense drugs. The present sequence was used to illustrate the invention.
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                                                                                                                                 Sequence 19 BP; 6 A; 5 C; 8 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                  Score 19; DB 1; Le
Pred. No. 1.5e+02;
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Disclosure; Page 100; 220pp; Japanese.
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100.0%;
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les 19; Conserv
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DT 26-AU
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431 CAGGACTCGGCTCACAT 449

19; Conservative

Matches

CAGGACTCGGCTCACACAT

13

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ADP87877 standard; DNA; 19 BP

RESULT 233
ADP87877/c
ID ADP878
XX
AC ADP878

ADP87877

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The present invention relates to novel 2',5'-oligoadenylic acid analogs and their pharmacologically- acceptable salts. The analogs are stable with superior antitumour and antiviral activity and so are useful in cancer or antiviral therapy e.g. as antisense drugs. The present sequence was used to illustrate the invention.
                                                                                                                                          1..5
/taga - //taga - OTHER
/mote= "2',4'-oxyethylene linkage in the sugar residues"
                                                                                                                                                                                                        /*tag= c
|mod_base= OTHER
|note= "2′,4′-oxyethylene linkage in the sugar residues"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Stable 2',5'-oligoadenylic acid analogs containing natural and modified nucleic acid units as well as unusual phosphate groups with excellent activity particularly antitumor, applicable in cancer or antiviral
         Cytostatic; virucide; 2'; 5'-oligoadenylic acid analog; antitumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2',5'-oligoadenylic acid analog related oligonucleotide #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4.2%; Score 19; DB 1; Length 19;
100.0%; Pred. No. 1.5e+02;
tive 0; Mismatches 0; Indels
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                                                                                                              /mod_base= OTHER
/note= "phosphorothioate backbone"
                                                                                                                                                                                                                                                                                       /note= "C-hydroxyethyl phosphate"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19 BP; 6 A; 5 C; 8 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 101; 220pp; Japanese.
                                                                         ocation/Qualifiers
                                                                                                                                                                                                                                                                           /mod base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                     19-NOV-2003; 2003WO-JP014748.
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Best Local Similarity 100...
Loca 19; Conservative
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                                                                                       ...19
                        antiviral; cancer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Koizumi M, Morita K;
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                                                                            Key
modified_base
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                                                 Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to novel 2',5'-oligoadenylic acid analogs and their pharmacologically- acceptable salts. The analogs are stable with superior antitumour and antiviral activity and so are useful in cancer or antiviral therapy e.g. as antisense drugs. The present sequence was used to illustrate the invention.
                                                                                                                                                                                 /*tag= a
/mod_base= OTHER
/note= "2',4'-oxyethylene linkage in the sugar residues"
                                                                                                                                                                                                                                       /*tag= c
/mod_base= OTHER
/note= "2',4'-oxyethylene linkage in the sugar residues"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Stable 2',5'-oligoadenylic acid analogs containing natural and modified nucleic acid units as well as unusual phosphate groups with excellent activity particularly antitumor, applicable in cancer or antiviral
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                             Cytostatic; virucide; 2'; 5'-oligoadenylic acid analog; antitumour; antiviral; cancer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2',5'-oligoadenylic acid analog related oligonucleotide #7.
            2',5'-oligoadenylic acid analog related oligonucleotide #6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match

4.2%; Score 19; DB 1; Length 19;
Best Local Similarity 100.0%; Pred. No. 1.5e+02;
Matches 19; Conservative 0; Mismatches 0; Indels
                                                                                                                                                         'note = "phosphorothioate backbone"
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                                                                                                                                                                                                                                                                                                                    /note= "C-hydroxyethyl phosphate"
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                                                                                                  Location/Qualifiers
1. 19
/*tag= d
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Koizumi M, Morita K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2004-460494/43.
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modified_base
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                                                                              Synthetic.
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AC ADP8
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Silverman RH,
         misc_feature
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03-FEB-1998;
                                     misc_feature
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                                                                                            29-OCT-1998
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                                                                                                                                                                                                                                                                                                                          The present invention relates to novel 2',5'-oligoadenylic acid analogs and their pharmacologically- acceptable salts. The analogs are stable with superior antitumour and antiviral activity and so are useful in cancer or antiviral therapy e.g. as antisense drugs. The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; telomerase; hTR; activator-antisense complex; malignant; enzyme; cleave; brain; tumour malignant glioma; breast tumour; renal cell cancer; melanoma; prostate cancer; leukemia; polycthemia vera; myeloma; sarcoma; Hoddkin's lymphoma; Waldenstrom's macroglobulinemia; heavy chain disease; carcinoma; chemotherapeutic; antisense; ss.
                                                                note- "2', 4' -oxyethylene linkage in the sugar residues"
                                                                                                                                                                                                                                                         Stable 2',5'-oligoadenylic acid analogs containing natural and modified nucleic acid units as well as unusual phosphate groups with excellent activity particularly antitumor, applicable in cancer or antiviral
                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oligo contained activator-antisense complex spA12-anti-hTR.
                                                                                                                                                                                                                                                                                                                                                                                                    4.2%; Score 19; DB 1; Length 19; 100.0%; Pred. No. 1.5e+02; ive 0; Mismatches 0; Indels
                                                                                          /mod_base= OTHER
/note= "C-hydroxyethyl phosphate"
                                                                                                                                                                                                                                                                                                                                                                                     Sequence 19 BP; 6 A; 5 C; 8 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                         Disclosure; Page 100; 220pp; Japanese.
                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                               /*tag= a
/mod base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                            76 GIGCITITIGCICCCGCGC 94
                                                                                                                                                                                                                                                                                                                                                                                                                                                       GIGCITITGCICCCCGCGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP.
                                                                                                                                                             19-NOV-2003; 2003WO-JP014748
                                                                                                                                                                                19-NOV-2002; 2002JP-00334731
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAV68470 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                  *tag= b
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nes 19; Conservative
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                                                                                                                                                                                                 (SANY ) SANKYO CO LTD
                                                                                                                                                                                                                    Koizumi M, Morita K;
                                                                                                                                                                                                                                       WPI; 2004-460494/43.
                                                                                                                        WO2004046161-A1
                                  modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Key
modified_base
                                                                         modified_base
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                                                                                                                                           03-JUN-2004
         Synthetic.
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                                                                                                                                                                                                                                                                                        therapy
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This represents an antisense oligonucleotide to the RNA component of human telomerase (hTR) comprised in the. The invention relates to an activator-antisense complex that comprises: (a) an antisense oligo, complementary to a 12-25 nucleotide portion of the RNA component of hTR, with a hydroxyl moiety at the first end; and (c) an activator of RNase L attached to the linker attached to the cirst end, and (c) an activator of RNase L attached to the linker. The activator-antisense complex may be used for inhibiting the growth of a telomerase expressing malignant cell or tumour. The complex is used to specifically cleave the ribonucleotide portion of a telomerase enzyme. The complex inhibits growth of telomerase expressing malignant cells from brain tumour malignant glioma, breast tumour renal cell cancer, melanoma, and prostate cancer. Many other malignancies and related disorders, may be treated including various acute and chronic leukemias, polychemia vera, Hodgkin's and non-Hodgkin's lymphomas, multiple cumours, including numerous sarcomas and carcinomas. The complex is preferrably administered in combination with a chemotherapeutic agent, becompain, actinomycin D, or neocarinostentin. The present sequence is an example of a modified antisense oligo comprised in an activator-antisense
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New RNase L activator-telomerase antisense complex - useful to inhibit
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/note= "nucleotide in reverse orientation 3'-5'"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      telomerase activity in telomerase-expressing malignancies.
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.00.0%; Pred. No. 1.6e+02;
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                                                                                                  /note= "3'-3' internucleotide linkage"
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"Sp5'A(2'p5'A)11-Bu2'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Li G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Pred. ...
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example; Page 53; 81pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (CLEV-) CLEVELAND CLINIC FOUND. (USSH ) US NAT INST OF HEALTH.
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98US-00018125.
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/*tag= '
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Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kondo S,
/note=
                                                                                                                                                                     /*tag=
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US09436060A-16.rng.sl

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This represents an antisense oligonucleotide to the RNA component of human telomerase (hTR) comprised in the. The invention relates to an activator-antisense complex that comprises: (a) an antisense oligo, activator-antisense complex that comprises: (a) an antisense oligo, complementary to a 12-25 nucleotide portion of the RNA component of hTR, with a hydroxyl moiety at the first end; and (b) a linker attached to the first end; and (c) an activator of RNAse L attached to the inher. The activator-antisense complex may be used for inhibiting the growth of a telomerase-expressing malignant cell or tumour. The complex is used to specifically cleave the ribonucleotide portion of a telomerase enzyme. The complex inhibits growth of telomerase expressing malignant cells from brain tumour malignant glitoma, breast tumour, renal cell cancer, brain tumour malignant glitoma, breast tumour, renal cell cancer, cancer. Many other malignancies and related disorders, may be treated including various acute and chronic leukemias, polycthemia vera, Hodgkin's and non-Hodgkin's lymphomas, multiple tumours, including numerous sarcomas and carcinomas. The complex is an expension of the complex is particularly either cisplatin, doxorubicin, michomy manners is an expensive the particularly either cisplatin, doxorubicin, michomy manners is an expense.
               melanoma; prostate cancer; leukemia; polycthemia vera; myeloma; sarcoma;
Hodgkin's lymphoma; Waldenstrom's macroglobulinemia; heavy chain disease;
carcinoma; chemotherapeutic; antisense; ss.
cleave; brain; tumour malignant glioma; breast tumour; renal cell cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            complex - useful to inhibit
                                                                                                                                                                                                       /note= "antisense oligo sequence claimed in claim 4"
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/note= "nucleotide in reverse orientation 3'-5'"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 telomerase activity in telomerase-expressing malignancies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Torrence PF;
                                                                                                                                                                                                                                                                                                    /*tag= c
/note= "3'-3' internucleotide linkage"
                                                                                                                                                                                                                                          /*tag= a
/note= "Sp5'A(2'p5'A)3-Bu2"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New RNase L activator-telomerase antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Silverman RH, Kondo S, Cowell JK,
                                                                                                                                                 Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLEV-) CLEVELAND CLINIC FOUND.
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                                                                                                              Homo sapiens
                                                                                                                                                                misc_feature
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                                                                                                                                                                                                                                                                                 misc_feature
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                                                                                         Synthetic.
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bleomycin, actinomycin D, or neocarzinostatin. The present sequence is an example of a modified antisense oligo comprised in an activator-antisense
                                                                                                                                                                   Sequence 20 BP; 6 A; 5 C; 8 G; 1 T; 0 U; 0 Other;
                                                                                      complex spA4-anti-hTR
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4.2%; Score 19; DB 1; Length 20;
100.0%; Pred. No. 1.6e+02;
tive 0; Mismatches 0; Indels
                                                                                            76 GIGCITITGCICCCGCGC 94
                          Local Similarity 100.
Les 19; Conservative
       Query Match
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Gaps

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The present sequence is that of a short interfering RNA, denoted hTR#2 sIRNA, a 19 bp sequence centred in the 26 bp loop in the longest single-stranded region of human telemerase RNA. The siRNA comprises sense and antisense nucleic acids that are complementary to each other except for 2 thymidine deoxynucleotides at both 3' overhangs. A nucleic acid that are complementary to each other except for 2 comprising the double-stranded region of this siRNA is specifically claimed. The invention relates to the discovery that double-stranded interfering RNAs which target telomerase RNA, or mRNA encoding telomerase interfering RNAs which target telomerase RNA, or mRNA encoding telomerase activity. Inhibition of telomerase in cancer cells leads to telomerase activity, end-to-end chromosomal fusion, and apoptosis. Interference of shortening, end-to-end chromosomal fusion, and apoptosis. Interference of contraception or sterilisation, for immunosuppression, for treatment of contraception or sterilisation, for immunosuppression, for treatment of yeast, parasite and fungal infections, and in antiinflammatory therapies. As telomerase is active in a limited number of cell types, e.g. tumour cells, germaine cells, certain stem cells of the haematopoietic system, T and B cells, sun-damaged skin, and proliferative cervix, most normal cells are not affected by telomerase RNA interference therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                              "double-stranded region, specifically referred to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /*tag= a
/label= Sticky_end
/label= sticky_end
fnote= "the 3' end of the complementary strand overhangs
the 5' end of this strand by the sequence 5'-TT-3'"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel nucleic acid encoding or comprising interfering RNAs which target telomerase RNA, useful for inhibiting telomerase activity for treating cancer, infertility and disorders of the immune system.
                                                                                                                                                                                                                                                         RNA interference; short interfering RNA; siRNA; telomerase; cancer; tumour; cytostatic; contraceptive; immunosuppressive; antiinfertility; fungicide; antiparasitic; antiinflammatory; human; gene therapy; ds.
                                                                                                                                                                                                                     Short interfering RNA hTR#2 siRNA, targets telomerase RNA.
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/label= Sticky_end
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19 GIGCITIIGCICCCCGCGC 1
                                                                                                 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22-OCT-2001; 2001US-0345326P.
20-FEB-2002; 2002US-0359196P.
22-MAY-2002; 2002US-0383195P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-OCT-2002; 2002WO-US033146.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   in Claim 11"
                                                                                                 ACC57540 standard; DNA; 21
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/*tag= b
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                                                                                                                                                                                                                                                                                                                                                                                       Key
misc_feature
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                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                         ACC57540;
                                                           RESULT 239
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Matches

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WO2003070742-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-AUG-2002;
05-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
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06-JUN-2002;
                                                                                                                                                                                                                                                                                  26-FEB-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
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                                                                                                                               Query Match
                                                                                                                                                                                                                       RESULT 241
ADF93815/c
8,566666666666
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence is that of a short interfering RNA, denoted hTR#1 siRNA, which targets the template region (see ACG57538) of human telomerase RNA. The siRNA comprises sense and antisense nucleic acids that are complementary to each other except for 2 thymidine deoxynucleotides at both 3' overhangs. A nucleic acid comprising the duble-stranded region of this siRNA is specifically claimed. The invention relates to the discovery that double-stranded interfering RNAs which target telomerase RNA, or mRNA encoding telomerase reverse
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                                                                                                                                                                                                                                                                                                                                                     /*tag= a
/label= Sticky_end
/label= "the 3' end of the complementary strand overhangs
the 5' end of this strand by the sequence 5'-TT-3'"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel nucleic acid encoding or comprising interfering RNAs which target telomerase RNA, useful for inhibiting telomerase activity for treating cancer, infertility and disorders of the immune system.
                                                                                                                                                                                                                      RNA interference; short interfering RNA; siRNA; telomerase; cancer; tumour; oytostatic; contraceptive; immunosuppressive; antiinfertility; fungicide; antiparasitic; antiinflammatory; human; gene therapy; DNA-RNA hybrid; ds.
                                                                                                                                                                                                                                                                                                              /*tag= b
/note= "double-stranded region, specifically referred
in Claim 10"
                                                 Gaps
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                                                                                                                                                                                                   Short interfering RNA hTR#1 siRNA, targets telomerase RNA
                             Score 19; DB 1; Length 21;
Pred. No. 1.7e+02;
                                                 0; Indels
          6 G; 5 T; 0 U; 0 Other;
                                  100.0%; Prea. ...
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/label= Sticky_end
                                                                   270 GGCTTCTCCGGAGGCACCC 288
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                                                                                     1 GGCTTCTCCGGAGGCACCC 19
                                                                                                                                        ACC57539 standard; DNA; 21 BP
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2002US-0383195P.
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                             4.2%;
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C;
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                                      Best Local Similarity 100.
Matches 19; Conservative
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          BP; 2 A;
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                                                                                                                                                                                                                                                                                                                                                                                            misc_feature
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          Sequence 21
                                                                                                                                                                               28-JUL-2003
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                                                                                                                                                            ACC57539
                             Query Match
                                                                                                                     RESULT 240
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transcriptase (TERT), are capable of inhibiting telomerase activity. Inhibition of telomerase in cancer cells leads to telomera shortening, end-to-end chromosomal fusion, and apoptosis. Interference of telomerase activity can also be used for treatment of infertility, for contraception or sterilisation, for immunosuppression, for treatment of yeast, parasite and fungal infections, and in antiinflammatory therapies. As telomerase is active in a limited number of cell types, e.g. tumour cells, germline cells, certain stem cells of the hematopojoitic system, T and B cells, sun-damaged skin, and proliferative cervix, most normal cells are not affected by telomerase RNA interference therapy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                Score 19; DB 1; I
Pred. No. 1.7e+02;
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2002US-0386782P.
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29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
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Matches 13; Conserv
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The invention relates to short interfering nucleic acids (sinA) which downregulate expression of the one or more telomerase genes by RNA interference. The sinAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and matisense regions, or alternatively are assembled from a sense oligonuclectide and an interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA, conjugates candor complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and cranded the conficions they may be conficions.
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                                                                                                                                                                                                                                                                                                                                                                                                              used for treating cancer, restenois, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siMs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nuclectide polymorphisms). The present sequence represents a chemically modified siRNA targeted to the human TERC mRNA
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telomerase gene useful e.g. for treatment and diagnosis of cancer.
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4.2%; Score 19; DD.
Best Local Similarity 100.0%; Pred. No. 1.7.
Best Local Similarity 0, Mismatches
                                               Example 3; SEQ ID NO 542; 145pp; English.
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interference. The siNAs may or may not comprise ribonucleotides and may be downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short hairpin RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised. Can contain deoxyribonucleotides, and can be chemically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cals, tissue caplants or the treatment of a variety of conditions. They may be transplants for the treatment of a variety of conditions. They may be protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, cannon in the sale ouseful for drug cannon in the case of the cannon cannon in the cannon cannon
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100.0%; Pred. No. 1.7e+02;
ive 0; Mismatches 0; Indels
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/note= "Ribothymidines. Also, the internucleotide linkage
is phosphorothioate"
/note= "Pyrimidine bases are 2'-deoxy-2'-fluoro and purines are Deoxy bases"
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                                                                                             4.2%; Score 19; DB 1; Length 21;
100.0%; Pred. No. 1.7e+02;
                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human TERC chemically modified siRNA, SEQ ID 539.
                                          Sequence 21 BP; 7 A; 2 C; 6 G; 2 T; 4 U; 0 Other;
                                                                                                                                             0; Mismatches
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                                                                                                                                                                                                   146 TTCCACCGTTCATTCTAGA 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       base= OTHER
                                                                                                                                                                                                                                TTCCACCGTTCATTCTAGA 1
                                                                                                                                                                                                                                                                                                                                                                                ADF93812 standard; RNA; 21 BP
                                                                                                                     100.08;
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2002US-0406784P.
2002US-0408378P.
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2002US-0386782P
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                                                                                                                                                19; Conservative
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                                                                                                                     Best Local Similarity
Matches 19; Conserv
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11-MAR-2002;
06-JUN-2002;
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transcript.
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                                                                                             Query Match
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regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised. Cerpressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants of organisms (e.g., by ex vivo gene therapy), or in grafts and transplants of organisms (e.g., by ex vivo gene therapy), or in grafts and transplants of organisms (e.g., by ex vivo gene therapy), or in grafts and transplant rejection, or autoimmune or age-related diseases, protozoal), transplant rejection, or autoimmune or age-related diseases. Ce.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents a chemically modified siRNA targeted to the human TERC mRNA
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Sequence 21 BP; 4 A; 7 C; 3 G; 2 T; 5 U; 0 Other;

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                           Gaps
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4.2%; Score 19; DB 1; Length 21; 73.7%; Pred. No. 1.7e+02; cive 5; Mismatches 0; Indels
                                                    148 CCACCGTTCATTCTAGAGC 166
                                                                   14; Conservative
                Best Local Similarity
    Query Match
                               Matches
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Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiartitc; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; DNA-RNA hybrid; ss. Human TERC chemically modified siRNA, SEQ ID 552. ADF93825 standard; RNA; 21 BP (first entry) 26-FEB-2004 Synthetic ADF93825;

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/mod_base= OTHER
/note= "Ribothymidines. Also, the internucleotide linkage
is phosphorothioate"
                                                                                                     /note= "Pyrimidine bases are 2'-deoxy-2'-fluoro"
                                Location/Qualifiers
                                                                                   /mod base= OTHER
                                                   ...21
*tag= b
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                                                                                                                                           *tag=
                                               modified_base
                                                                                                                          modified base
Homo sapiens.
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WO2003070742-A1

20-FEB-2002; 2002US-0358580P.

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11-FEB-2003; 2003WO-US004088
28-AUG-2003.
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense or egions, or alternatively are assembled from a sense onigonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short can contain deoxyribonucleotides, and can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised, can complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue caphants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, respenses infectious diseases (specifically protococal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, ALDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, lupus erythematosus, ALDS, macular degeneration, cancering, pharmacogenomics, studying gene function, and gene corpused a chemically modified siRNA targeted to the human TERC mRNA
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                                                                                          05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
15-JAN-2003; 2003US-0440129P.
                       06-JUN-2002; 2002US-0386782P.
17-JUL-2002; 2002US-0396600P.
29-AUG-2002; 2002US-0406784P.
11-MAR-2002; 2002US-0363124P.
                                                                                                                                                                                                 (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                             Beigelman L;
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0; Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering nucleic acid; siNA; short siRNA; double-stranded RNA; micro-RNA; mixNA; short haripin RNA; shrNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; Gaps .; 0 4.2%; Score 19; DB 1; Length 21; 100.0%; Pred. No. 1.7e+02; ive 0; Mismatches 0; Indels Human TERC chemically modified siRNA, SEQ ID 544. 300 GAAGAGTTGGGCTCTGTCA 318 19 GAAGAGTTGGGCTCTGTCA 1 BP ADF93817 standard; RNA; 21 26-FEB-2004 (first entry) Query Match
Best Local Similarity 100.
Matches 19; Conservative ADF93817; RESULT 246 ADF93817 ò ద

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ADF93832/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonuclectide and antisense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short harpin RNA (shRNA). The siNAs can be unmodified or chemically modified, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and vectors that express since in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, c. e.g. multiple sclerosis, lupus exythematosus, ADS, macular degeneration, explants or organisms therapeutic target identification and validation, and send multiple sclerosis, lupus expthematosus, ADS, macular degeneration, and send multiple sclerosis, lupus expthematosis, and send function, and send manning of aincle manning send expenses.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mapping (e.g., of single nucleotide polymorphisms). The present sequence represents a chemically modified siRNA targeted to the human TERC mRNA
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100.0%; Pred. No. 1.7e+02;
ive 0; Mismatches 0; Indels
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                                                                                                                             /*tag= a
/mod_base= OTHER
/note= "Ribothymidines"
                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                             2002US-0396600P.
2002US-0406784P.
2002US-0408378P.
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2002US-0386782P
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DNA-RNA hybrid; ss.
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                                                                                                             modified base
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29-AUG-2002;
                                                          Homo sapiens
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                                       Synthetic
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antiennse regions, or alternatively are assembled from a sense oligonucleotide and an antistense oligonucleotide. Specifically, the siNAs include short interfering RNA (siNAs), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siNNA), double-stranded RNA, micro-RNA (miRNA) and short can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised, me invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are
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/mod_base= OTHER
/note= "Ribothymidines. Also, the internucleotide linkage
is phosphorothloate"
                                                                                                                            Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiartitic; antiinflammatory; gene therapy; telonmerase; human; terc; RNA interference; short interfering nucleic acid; siNN; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERT; DNA-RNA hybrid; ss.
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/note= "Pyrimidine bases are 2'-deoxy-2'-fluoro and
purines are Deoxy bases"
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                                                                                              Human TERC chemically modified siRNA, SEQ ID 559.
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; 2002US-0408378P.
; 2002US-0409293P.
; 2003US-0440129P.
ADF93832 standard; RNA; 21 BP
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2002US-0386782P.
2002US-0396600P.
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                                                               (first entry)
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modified_base
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05-SEP-2002;
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                               ADF93832;
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Gaps

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300 GAAGAGTIGGGCTCTGTCA 318

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Conservative

Local Similarity les 19; Conserv

Best Loca Matches

19 GAAGAGTTGGGCTCTGTCA 1

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used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenceis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple solarons, lupus erythematocusus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siMAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents a chemically modified siRNA targeted to the human TERC mRNA
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100.0%; Pred. No. 1.7e+02;
iive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 5 A; 3 C; 7 G; 2 T; 4 U; 0 Other;
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/mod_base= OTHER
/note= "Ribothymidines"
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11-MAK-2002; 2002US-0363124P.

06-JUN-2002; 2002US-0386782P.

17-JUL-2002; 2002US-0396600P.

29-AUG-2002; 2002US-0496784P.

05-SEP-2002; 2002US-040878P.

09-SEP-2002; 2002US-0409293P.
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nes 19; Conservative
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modified_base
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Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            L5-JAN-2003;
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                                                                                                                                                                                                                                                                                                                                                                               transcript
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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WPI; 2003-689777/65.

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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an internee oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised. Cexpressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro of synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; double-stranded RNA; micro-RNA; minA; short halfpin RNA; shrNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents a chemically modified siRNA targeted to the human TERC mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Gaps
                 New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4.2%; Score 19; DB 1; Length 21;
53.2%; Pred. No. 1.7e+02;
ve 7; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 4 A; 6 C; 2 G; 2 T; 7 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human TERC chemically modified siRNA, SEQ ID 540.
                                                                                         Example 3; SEQ ID NO 538; 145pp; English.
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/mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 63.2%;
Marches 12; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA-RNA hybrid; ss.
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ranscript.
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transcript
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downregulate expression of the one or more telemerase genes by NNA interference. The siNAs may or may not comprise sense and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the siNAs include short in the ferring RNA (siRNA), double-stranded RNA, micro-RNA (mixRNA) and short can contain deoxyribonucleotides, and can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised. The invention also crelates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express sink. The siNAs are used to modulate expression of the telomerase genes in cells, tissue caplants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be conditions. The transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosus, AlDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, disagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene constants a chemically modified siRNA targeted to the human TERC mRNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc;
                                                                                                                                                                                                                                                                                                              The invention relates to short interfering nucleic acids (siNA) which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                            New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 19; DB 1; Length 21; Pred. No. 1.7e+02; 5; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 4 A; 3 C; 7 G; 2 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human TERC chemically modified siRNA, SEQ ID 560.
                                                                                                                                                                                                                                                                                   Example 3; SEQ ID NO 540; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     300 GAAGAGTTGGGCTCTGTCA 318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 GAAGAGUUGGCUCUGUCA 19
                                               20-FEB-2002; 2002US-0358580P.
11-MAX-2002; 2002US-0363124P.
06-UJN-2002; 2002US-0386782P.
17-UJL-2002; 2002US-0396600P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0409378P.
09-SEP-2002; 2002US-0409293P.
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                                                                                                                                         15-JAN-2003; 2003US-0440129P
                        11-FEB-2003; 2003WO-US004088
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                                                                                                                                                                  (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                          Beigelman L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                     WPI; 2003-689777/65.
                                                                                                                                                                                           Mcswiggen J,
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28-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    transcript.
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The invention relates to short interfering nucleic acids (siNh) which downregulate expression of the one or more telomerase genes by RNA interference. The siNhs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and mantisense regions, or alternatively are assembled from a sense oligonuclectide and antisense oligonuclectide. Specifically, the siNhs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNhs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised. Can contain deoxyribonuclectides, and can be chemically synthesised. Can complexes of siNA; and vectors that express siNA. The siNhs are and/or complexes of siNA; and vectors that express siNA. The siNhs are used to modulate expression of the telomerase genes in cells, tissue cand/or complexes of siNA; and vectors that express siNA. The siNhs are used to modulate expression of the telomerase genes in cells, tissue used to modulate expression of the telomerase genes in cells, transplant circumant of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically cransplants for the treatment of a variety of conditions. They may be used for transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus erythematosuus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNhs are also useful for drug screening, diagnosis, therapeleutic target identification and validation, genetic engineering, pharmacogenomics, studding gene function, and gene mapping (e.g., of single silknA targeted to the human TERC mRNA transparents a chemically modified siRNA targeted to the human TERC mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "Ribothywidines. Also, the internucleotide linkage is phosphorothioate"
                                                miRNA;
RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA short halrpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT; DNA-RNA hybrid; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note= "Pyrimidine bases are 2'-deoxy-2'-fluoro and purines are Deoxy bases"
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/mod_base= OTHER
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2002US-0386782P.
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2002US-0406784P.
2002US-0408378P.
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*tag=
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29-AUG-2002;
05-SEP-2002;
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06-JUN-2002;
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                                          Length 21;
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  BP; 5 A; 7 C; 3 G; 2 T; 4 U; 0 Other;
                                        4.2%; Score 19; DB 1; Le
100.0%; Pred. No. 1.7e+02;
:ive 0; Mismatches 0;
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                                                                                                                                  300 GAAGAGTTGGGCTCTGTCA 318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   base= OTHER
                                                                                                                                                        GAAGAGTTGGGCTCTGTCA 1
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2002US-0363124P.
2002US-0386782P.
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29-MUG-2002, 2002US-0406784P-
05-SEP-2002, 2002US-0408378P-
09-SEP-2002, 2002US-0409293P-
15-JAN-2003, 2003US-0440129P-
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                                                                                                                                                                                                                                                                                   ADF93816 standard; RNA; 21
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                                                                 Local Similarity 100, nes 19; Conservative
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modified base
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      Sequence 21
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                                                                                                                                                                                                                                                                                                                              ADF93816;
                                                                                                                                                                            19
                                               Query Match
                                                                                                                                                                                                                                           RESULT 251
                                                                                           Matches
                                                                                                                                                                                                                                                             ADP 93816/

AC ADP 93

AC ADP 93
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hairpin RNA (shRNA). The sinAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of sinA, conjugates and/or complexes of sinA, and vectors that express sinA. The sinAs are used to modulate expression of the telomerase genes in cells, tissue can organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, protozoal), transplant rejection, or autoimmune or age-related diseases, sin ulcers and rheumatoid arthritis. The sinAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence mapping (e.g., of single mucleotide polymorphisms). The present sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthitic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; abhort hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /mod_base= OTHER
/note= "Pyrimidine bases are 2'-deoxy-2'-fluoro"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21 BP; 5 A; 3 C; 7 G; 2 T; 4 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4.2%; Score 19; DB 1; Le
100.0%; Pred. No. 1.7e+02;
tive 0; Mismatches 0;
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base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             148 CCACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19 CCACCGTTCATTCTAGAGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2002US-0358580P.
2002US-0363124P.
2002US-0386782P.
2002US-0396600P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ρ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity 100.
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20. .21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA-RNA hybrid; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO2003070742-A1
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modified_base
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06-JUN-2002;
17-JUL-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          modified base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-FEB-2002;
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonuclectide and interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonuclectides, and can be chemically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA. The siNAs are used to modilate expression of the telomerase genes in cells, tissue and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modilate expression of the telomerase genes in cells, tissue caplants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants of the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically crocoal), transplant rejection, or autoimmune or age-related diseases, e.g. multiple sclerosis, lupus expthematosus, AlbS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single mucleotide polymorphisms). The present sequence mapping (e.g., of single mucleotide polymorphisms). The present sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         double-stranded short interfering nucleic acid; siNA; antiarteriosclerotic; neuroprotective; noctropic; antiparkinsonian; anticonvulsant; pulmonary disease; restenosis; atherosclerosis; allaheimer's; Parkinson's; epilepsy; dementia; huntington's; amyotrophic lateral sclerosis; gene therapy; ss; DNA-RNA hybrid; hTR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                  New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4.2%; Score 19; DB 1; Length 21;
100.0%; Pred. No. 1.7e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 7 A; 2 C; 6 G; 2 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sina DNA-RNA hybrid - SEQ ID 608.
                                                                                                                                                                                                                                                                                                    Example 3; SEQ ID NO 550; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     146 TICCACCGITCATICIAGA 164
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29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409291P.
15-JAN-2003; 2003US-04401291P.
                                                                                                        (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                  Beigelman L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
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                                                                                                                                                                                          WPI; 2003-689777/65.
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nes 19; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          402003074654-A2
                                                                                                                                                  Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   hTR-targeted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unidentified
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The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of the invention demonstrates antiarteriosclerotic, neuroprotective, conotropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition target gene and therefore in the treatment of any disease or condition target gene and target gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary diseases such as restenosis, atherosclerosis, Alzheimer's disease, Parkinson's disease, epilepsy, dementia, huntington's disease or myctrophic lateral sclerosis. Furthermore, the siNA may be utilised for gene therapy applications. The current sequence is that of the siNA DNA-RNA hybrid of the invention.
                                                                                                                                                                                                                                                                                                                                         New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antiarteriosclerotic; neuroprotective; nootropic; antiparkinsonian; anticonvulaent; pulmonary disease; restenoslas; atherosclerosis; Alzheimer's; Parkinson's; epislepsy; dementia; huntington's; amyotrophic lateral sclerosis; gene therapy; ss; DNA-RNA hybrid; hTR.
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                                                                                                                                                                                                                                                      Fosnaugh K;
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Pred. No. 1.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
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                                                                                                                                                                                                                                                      Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 21 BP; 7 A; 2 C; 6 G; 2 T; 4 U; 0 Other;
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                                                                                                                                                                                                                                                      Chowrira B,
                                                                                                                                                                                                                                                                         Thompson J;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TICCACCGTICATICIAGA 1
                                                                                                                                                                                                                  (SIRN-) SIRNA THERAPEUTICS INC
                                                                                    2002US-0363124P.
2002US-0386782P.
2002US-0406784P.
2002US-0408378P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 100.0%; es 19; Conservation
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                                                                                                                                                          2002US-0409293P
                                                                                                                                                                               15-JAN-2003; 2003US-0440129P.
                                 20-FEB-2003; 2003WO-US005028
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                                                                                                                                                                                                                                                      Beigelman
                                                                                                                                                                                                                                     Mcswiggen J, Belyerman S. Usman N,
                                                                                                                                                                                                                                                                                                           WPI; 2003-731676/69.
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                                                                                                                      29-AUG-2002;
05-SEP-2002;
09-SEP-2002;
                                                                                                           36-JUN-2002;
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                                                                                      11-MAR-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                   activity.
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WO2003074654-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (siNA) molecule that down-regulates expression of an endogenous mammalian terget gene comparising one or more chemical modifications and each strand of the double-stranded sink comprises about 21 nuclectides. The siNA of the invention demonstrates antiarteriosclerotic, neuroprotective,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a double-stranded short interfering nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nootropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regularing the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a call, tissue or organism. The disease or condition may include pulmonary diseases such as restenois, athereiser's disease, Parkinson's disease, epilepsy, dementia, huntington's disease or amyorrophic lateral solerosis. Furthermore, the SINA may be utilized for
                                                                                                                                                                                                                                                                                                                                                                                                                            New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              gene therapy applications. The current sequence is that of the siNA DNA-RNA hybrid of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    double-stranded short interfering nucleic acid; siNA; antiarteriosclerotic; neuroprotective; noctropic; antiparkinsonian; anticonvulsant; pulmonary disease; restenosis; atherosclerosis; alzheimer's; Parkinson's; epilepsy; dementia; huntington's; amyotrophic lateral sclerosis; gene therapy; ss; DNA-RNA hybrid; hTR.
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0
                                                                                                                                                                                                                                                                                                                          Fosnaugh K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4.2%; Score 19; DB 1; Length 21; 100.0%; Pred. No. 1.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                          Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 5 A; 3 C; 7 G; 2 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            hTR-targeted sina DNA-RNA hybrid - SEQ ID 610.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Pred. w.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 24; SEQ ID NO 609; 593pp; English.
                                                                                                                                                                                                                                                                                                                          Chowrira B,
                                                                                                                                                                                                                                                                                                                 Beigelman L, Chowr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               148 CCACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ccaccerrcarrcragage 1
                                                                                                                                             2002US-0386782P.
2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
                                                                                                                                                                                                                                                                               (SIRN-) SIRNA THERAPEUTICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADG30044 standard; RNA; 21 BP
                                                                                                       2002US-0358580P.
2002US-0363124P.
                                                                                                                                                                                                                                       2003US-0440129P
                                                              20-FEB-2003; 2003WO-US005028
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19; Conservative
                                                                                                                                                                                                                                                                                                     Mcswiggen J, Belyer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-731676/69.
                                                                                                                                               06-JUN-2002;
29-AUG-2002;
05-SEP-2002;
09-SEP-2002;
                                                                                                                                                                                                                                       15-JAN-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unidentified
                                                                                                                             11-MAR-2002;
                      12-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  activity.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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NAME OF THE PROPERTY OF THE PR
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The invention relates to a double-stranded short interfering nucleic acid (siN3) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 2; nucleotides. The siNA of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            the invention demonstrates antiarteriosclerotic, neuroprotective, nootropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treament of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary diseases such as restenosis, atherosclerosis, Albreimer's disease, Parkinson's disease, epilepsy, dementia, huntington's disease or amyotrophic lateral sclerosis. Purthermore, the sinA may be utilised for gene therapy applications. The current sequence is that of the siNA DNA-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RNA interference; short interfering RNA; siRNA; telomerase; cancer; tumour; cytostatic; contraceptive; immunosuppressive; antiinfertility; fungicide; antiparasitic; antiinflammatory; human; gene therapy; DNA-RNA hybrid; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                Fosnaugh K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Short interfering RNA hTR#2 siRNA, targets telomerase RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 19; DB 1; Length 21; Pred. No. 1.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
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100.0%; Pred. No. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 24; SEQ ID NO 610; 593pp; English.
                                                                                                                                                                                                                                                                                                                                                                                         Beigelman L, Chowri
sman N, Thompson J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 300 GAAGAGTTGGGCTCTGTCA 318
                                                                                         11-MAR-2002; 2002US-0363124P.
06-JUN-2002; 2002US-0386782P.
29-AGG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2003; 2002US-040923P.
15-JAN-2003; 2003US-0440129P.
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                                                                                                                                                                                                                                                                                                                              (SIRN-) SIRNA THERAPEUTICS INC
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2003WO-US005028
                                                              2002US-0358580P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RNA hybrid of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-AUG-2003 (first entry)
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Best Local Similarity 100.
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                               Usman N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-731676/69.
20-FEB-2003;
                                                                                                                                                                                                                                                                                                                                                                                            Mcswiggen J,
                                                              20-FEB-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                               Jamison S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
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Location/Oualifiers

/*tag= b /note= "double-stranded region, specifically referred to

in Claim 10"

misc_feature

Location/Qualifiers

...19 *tag=

misc_feature

Synthetic.

RNA interference; short interfering RNA; siRNA; telomerase; cancer; tumour; cytosteatic; contraceptive; immunosuppressive; antinfertility; fungicide; antiparasitic; antinflammatory; human; gene therapy; DNA-RNA hybrid; ss.

Short interfering RNA hTR#1 siRNA, targets telomerase RNA.

overhangs

//tag= a //label= Sticky_end //note= "the 3' end of the complementary strand ove the 5' end of this strand by the sequence 5'-TT-3'

/*tag= c /label= Sticky_end

WO2003035667-A2

31-MAY-2003

. 21

misc_feature

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The present sequence is that of a short interfering RNA, denoted hTR#2 siRNA, a 19 bp sequence centred in the 26 bp loop in the longest single-stranded region of human tellomerase RNA. The siRNA comprises sense and antisense nucleic acids that are complementary to each other except for 2 thymidine deoxynucleotides at both 3' overhangs, A nucleic acid that size at both 3' overhangs, A nucleic acid comprising the double-stranded region of this siRNA is specifically claimed. The invention relates to the discovery that double-stranded interfering RNAs which target telomerase RNA, or mRNA encoding telomerase cremescriptase (TRRT), are capable of inhibiting telomerase activity. Inhibition of telomerase in cancer cells leads to telomere shortening, end-to-end chromosomal fusion, and apoptosis. Interference of telomerase activity can also be used for treatment of infertility, for contraception or sterilisation, for immunosuppression, for treatment of yeast, parasite and fungal infections, and in antiinflammatory therapies. As telomerase is active in a limited number of cell types, e.g. tumour cells, germline cells, certain stem cells of the haematopoietic system, T and B calls, sun-damaged skin, and proliferative cervix, most normal cells are not affected by telomerase RNA interference therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel double-stranded short interfering RNA having sense and antisense unucleic acids which are complementary to each other and to target nucleic acid e.g., telomerase RNA or mRNA encoding telomerase reverse transcriptase.
                 /*tag= b
/note= "double-stranded region, specifically referred to
                                                                                           /*tag= a
/label= Sticky_end
/note= "the 3' end of the complementary strand overhangs
the 5' end of this strand by the sequence 5'-TT-3'"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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100.0%; Pred. No. 1.7e+02;
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                                                                                                                                                                                        /*tag= c
/label= Sticky end
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 11; Page 28; 37pp; English
                                                                                                                                                                                                                                                                                                                                                           22-OCT-2001; 2001US-0345326P.
20-FEB-2002; 2002US-0359196P.
22-MAY-2002; 2002US-0383195P.
                                                                                                                                                                                                                                                                                                                       16-OCT-2002; 2002WO-US033065
                                                     in Claim 11"
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                                                                                                                                                                                                                                                                                                                                                                                                                                      (UYRP ) UNIV ROCHESTER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-403336/38
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                                                                                                                                                                                                                                                                                                                                                         22-OCT-2001;
                                                                                                                                                                misc_feature
 misc_feature
                                                                           misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rowley PT;
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22-OCT-2001; 2001US-0345326P. 20-FEB-2002; 2002US-0359196P. 22-MAY-2002; 2002US-0383195P.

(UYRP) UNIV ROCHESTER

WPI; 2003-403336/38.

PT;

Rowley

transcriptase.

16-OCT-2002; 2002WO-US033065.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence is that of a short interfering RNA, denoted hTR#1 siRNA, which targets the template region (see ACC57526) of human telomerase RNA. The siRNA comprises sense and antisense nucleic acids that are complementary to each other except for 2 thymidine deoxynucleotides at both 3' overhangs. A nucleic acid comprising the double-stranded region of this siRNA is specifically claimed. The invention relates to the discovery that double-stranded interfering RNAs which target telomerase RNA, or mRNA encoding telomerase activity. Inhibition of telomerase in cancer cells leads to telomerase activity. Inhibition of telomerase activity end-to-end chromosomal fusion, and apoptosis. Interference of telomerase activity can also be used for treatment of infertility, for contraception or sterilisation, for immunosuppression, for treatment of yeast, paragite
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 and fungal infections, and in antiinflammatory therapies. As telomerase is active in a limited number of cell types, e.g. tumour cells, germline cells, certain stem cells of the haematopoietic system, T and B cells, sun-damaged skin, and proliferative cervix, most normal cells are not
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58.4%; Pred. No. 1.7e+02;
tve 6; Mismatches 0; Indelb
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Best Local Similarity 68.4°
Matches 13, Conservative
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270 GGCTTCTCCGGAGGCACCC 288

1 GGCTTCTCCGGAGGCACCC 19

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ACC58027 standard; DNA; 21 BP

RESULT 257

ACC58027

(first entry)

11-AUG-2003

ACC58027;

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Novel double-stranded short interfering RNA having sense and antisense nucleic acids which are complementary to each other and to target nucleic acid e.g., telomerase RNA or mRNA encoding telomerase reverse

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modified base
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                                                                                                                       Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; siRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacognomics; gene function analysis; gene mapping; TERC; TERT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to short interfering nucleic acids (siNA) which
                                                                                                                                                                                                                                                                              /note= "Pyrimidine bases are 2'-deoxy-2'-fluoro and purines are Deoxy bases"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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/mod_base= OTHER
/note= "Inverted deoxy abasic nucleotide"
                                                                                                                                                                                                                                                                                                                          /note= "Inverted deoxy abasic nucleotide"
21. .22
                                                                                                         Human TERC chemically modified siRNA, SEQ ID 556.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 3; SEQ ID NO 556; 145pp; English
                                                                                                                                                                                                                                                                                                                                                               /note= "Ribothymidines"
                                                                                                                                                                                                                                            Location/Qualifiers
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/mod_base= OTHER
                                                                                                                                                                                                                                                                      /mod_base= OTHER
                                                                                                                                                                                                                                                                                                                   base= OTHER
::|:|:|||||:|||| ungucuaaccuaacugag 19
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11-MAR-2002, 2002US-0363144P.
06-JUN-2002, 2002US-03660PB.
17-JUL-2002, 2002US-036600P.
29-AUG-2002, 2002US-0406784P.
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2002US-0409293P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      11-FEB-2003; 2003WO-US004088
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                                                     ADF93829 standard; RNA; 23
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                                                                                        (first entry)
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                                                                                                                                                                                                  DNA-RNA hybrid; ss.
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                                                                                                                                                                                                                                                      modified base
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                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-SEP-2002;
                                                                                         26-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                      28-AUG-2003
                                                                                                                                                                                                                    Synthetic
                                                                       ADF93829;
                                   RESULT 25
ADF93829
                                                      合
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cc downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise stabonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siNRA), double-stranded RNA, micro-RNA (miKNA) and short haippin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also crelates to kits for the in vitro or in vivo delivery of siNA, conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treatgains (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be conditible sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, subdying gene function, and gene mapping (e.g., of single mucleotide polymorphisms). The present sequence represents a chemically modified siRNA targeted to the human TERC mRNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /mod_base= OTHER
/note= "Pyrimidine bases are 2'-deoxy-2'-fluoro and
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/note= "Inverted deoxy abasic nucleotide"
21. .22
/*tag= c
/mod_base= OTHER
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/note= "Ribothymidines"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 4.2%;
Best Local Similarity 73.7%;
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADF93828 standard; RNA; 23
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Human TERC chemically modified siRNA, SEQ ID 554.

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The invention relates to short interfering nucleic acids (sinh) which downregulate expression of the one or more telomerase genes by RNA interference. The sinham may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense oligonucleotide. Specifically, the siNhs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNhs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNhs are used to modulate expression of the telomerase genes in cells, tissue used to modulate expression of the telomerase genes in cells, tissue transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically proteors and relating delation) or autoimmune or age-related diseases, proteors and relating and relating dath are also useful for drug skin ulcers and rheumatoid arthritis. The sinha are also useful for drug cancer, cannot an analysic of condition and validation, cancer in calls.
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/*tag= d
/mod_base= OTHER
/note= "Inverted deoxy abasic nucleotide"
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                                                                                                                                                                                                                                                                        2002US-0396600P.
2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
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2002US-0363124P.
2002US-0386782P.
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                                                                                                                                                                                                                                                                                                                                                                                                       (RIBO-) RIBOZYME PHARM INC.
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06-JUN-2002;
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antienne regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, can contain a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are
                                        Cytostatic; vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antiulcer; antirheumatic; antiartitic; antiinflammatory; gene therapy; delomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERT; DNA-RNA hybrid; ss.
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/mod_base= OTHER
/note= "Inverted deoxy abasic nucleotide"
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/note= "Inverted deoxy abasic nucleotide"
21. .22
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/note= "Ribothymidines"
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/mod_base= OTHER
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2002US-0408378P.
2002US-0409293P.
2003US-0440129P.
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2002US-0386782P.
2002US-0396600P.
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Gaps

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4.2%; Score 19; DB 1; Length 23; 73.7%; Pred. No. 1.9e+02; cive 5; Mismatches 0; Indels

148 CCACCGTTCATTCTAGAGC 166

Local Similarity 73.7

Matches

Query Match

CCACCGUUCAUUCUAGAGC 20

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ADF93827 standard; RNA; 23 BP

26-FEB-2004 (first entry)

ADF93827;

RESULT 260
ADF93827
ID ADF9382
XX
AC ADF9382
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DT 26-FEB-

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explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, es, multiple solarosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siMAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents a chemically modified siRNA targeted to the human TERC mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cytostatic, vasotropic; protozoacide; immunosuppressive; dermatological; neuroprotective; anti-HIV; ophthalmological; antialcer; antirheumatic; antiarthritic; antiinflammatory; gene therapy; telomerase; human; terc; RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; TERC; TERT;
to modulate expression of the telomerase genes in cells, tissue
                                                                                                                                                                                                                                                                                      Gaps
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/note= "Inverted deoxy abasic nucleotide"
21. .22
                                                                                                                                                                                                                 Sequence 23 BP; 4 A; 6 C; 2 G; 2 T; 7 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human TERC chemically modified siRNA, SEQ ID 546.
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/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                       146 TICCACCGIICATICIAGA 164
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11-MAR-2002; 2002US-0363124P
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les 12; Conserv
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                                                                                                                                                                                                                                                    Query Match
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and matisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short interfering RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain decoxyribonucleotides, and can be chemically synthesised. Can contain decoxyribonucleotides, and can be chemically synthesised, can contain decoxyribonucleotides, and can be chemically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or age-related diseases, construction and shear also useful for drug screening, diagnosis, therapeutic target identification and validation, screening, cancer to redement of a varietic dentification and validation, and screening, cancer the sample of the carget identification.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 23 BP; 4 A; 6 C; 2 G; 2 T; 7 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human TERC chemically modified siRNA, SEQ ID 547.
                                                                                                                                                                                                                                                                                                                                                                                 Example 3; SEQ ID NO 546; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                146 TTCCACCGTTCATTCTAGA 164
                   2002US-0396600P.
2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
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2002US-0386782P
                                                                                                                15-JAN-2003; 2003US-0440129P
                                                                                                                                                               RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                               Beigelman L;
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Matches 12; Conservative
                                                                                                                                                                                                                                                            WPI; 2003-689777/65.
                                                                                                                                                                                                                 Mcswiggen J,
                                            29-AUG-2002;
                                                                05-SEP-2002;
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The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise ribonuclectides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonuclectide and antisense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically synthesised, can contain deoxyribonuclectides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue can for an expression of the telomerase genes in cells, tissue transplants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restenosis, infectious diseases (specifically proposed), transplant rejection, or autoimmune or age-related diseases, conditions and the conditions and the conditions are served.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         e.g. multiple sclerosis, lupus erythematosus, AIDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents a chemically modified siRNA targeted to the human TERC mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
                                                                                                                                               'note= "Pyrimidine bases are 2'-deoxy-2'-fluoro"
                                                                                                                                                                                                                                                                                                                                    /*tag= d
/mod_base= OTHER
/note= "Inverted deoxy abasic nucleotide"
                                                                                                                                                                                    /*tag= a
/mod_base= OTHER
/mode= "Inverted deoxy abasic nucleotide"
21. .22
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                                                                                                                                                                                                                                                                                                 'note= "Ribothymidines"
                                                                         Location/Qualifiers
                                                                                                                            base= OTHER
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mod base= OTHER
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2002US-0363124P.
2002US-0386782P.
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2002US-0406784P.
2002US-0408378P.
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15-JAN-2003; 2003US-0440129P
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                                                                                                          *tag=
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29-AUG-2002; 2
05-SEP-2002; 2
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modified_base
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06-JUN-2002;
                                    Homo sapiens
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                   Synthetic
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                                                Gарв
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                     4.2%; Score 19; DB 1; Length 23; 73.7%; Pred. No. 1.9e+02; tive 5; Mismatches 0; Indel8
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/mod_base= OTHER
/not== "Inverted deoxy abasic nucleotide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "Inverted deoxy abasic nucleotide"
Sequence 23 BP; 4 A; 7 C; 3 G; 2 T; 5 U; 2 Other;
                                                                                                                                                                                                                                     Human TERC chemically modified BiRNA, SEQ ID 548.
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                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                        148 CCACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                              /mod_base= OTHER
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/*tag= c
/mod_bage= OTHER
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                                                                                       CCACCGUUCAUUCUAGAGC 20
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2002US-0386782P.
2002US-039660P.
2002US-0406378P.
2002US-0408378P.
                                                                                                                                                             ADF93821 standard; RNA; 23 BP
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                        Query Match
Best Local Similarity 73.73
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                    modified base
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C downregulate expression of the one or more telomerase genes by RNA interference. The siNAs may or may not comprise response by RNA interference. The sinAs may or may not comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense regions, or alternatively are assembled from a sense oligonucleotide and antisense content and sense oligonucleotide. Specifically, the siNAs include short nation, and sense oligonucleotides, and can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised. The invention also can contain deoxyribonucleotides, and can be chemically synthesised. Cc can contain deoxyribonucleotides, and can be chemically synthesised, correlates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the telomerase genes in cells, tissue cerplants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating cancer, restencsis, infectious diseases (specifically protozoal), transplant rejection, or autoimmune or ager-elated diseases, e.g. multiple sclerosis, lupus erythematosus, ALDS, macular degeneration, skin ulcers and rheumatoid arthritis. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene correction, and gene correction.

C proposally reserved to single nucleotide polymorphisms). The present sequence correction.
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                                                                                                           The invention relates to short interfering nucleic acids (siNA) which
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New short interfering nucleic acid downregulates expression of the telomerase gene useful e.g. for treatment and diagnosis of cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    double-stranded short interfering nucleic acid; siNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 23 BP; 4 A; 3 C; 7 G; 2 T; 5 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hTR-targeted sina DNA-RNA hybrid - SEQ ID 605.
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                                                                Example 3; SEQ ID NO 548; 145pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              300 GAAGAGTTGGGCTCTGTCA 318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 GAAGAGUUGGGCUCUGUCA 20
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2002US-0386782P.
2002US-0406784P.
2002US-0408378P.
2002US-040929P.
2003US-0440129P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
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29-AUG-2002;
05-SEP-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        transcript
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The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of the invention demonstrates antiarteriosaclerotic, neuroprotective, nototropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition tissue or corganism. The disease or condition may include pulmonary diseases such as restenosis, atherosclerosis, Alzheimer's disease or Parkinson's disease, epilepsy, dementia, huntington's disease or manyotrophic lateral sclerosis. Furthermore, the siNA may be utilised for many thereary and include manyotrophic lateral sclerosis. Furthermore, the siNA may be utilised for many thereary and include manyotrophic actions and proper and pro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              gene therapy applications. The current sequence is that of the siNA DNA-RNA hybrid of the invention.
                                                                                                                                                                                                 New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or
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                                                                               Pavco P, Fosnaugh K;
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                                                                                                                                                                                                                                                                                                                                   Example 24; SEQ ID NO 605; 593pp; English.
                                                                            Chowrira B,
                                                                                                          Thompson J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             148 CCACCGITCATICIAGAGC 166
                       (SIRN-) SIRNA THERAPEUTICS INC.
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2002US-0408378P.
2002US-0409293P.
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2002US-0386782P.
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                                                                               Beigelman L,
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                                                     Mcswiggen J, Belyer...
G Usman N,
                                                                                                                                                       WPI; 2003-731676/69
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11-MAR-2002; 2
06-JUN-2002; 2
29-AUG-2002; 2
05-SEP-2002; 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12-SEP-2003.
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                                                                                                                                                                                                                                                                                 activity.
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RESULT 267
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à
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                셤
                                                                                                                                                                                          The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian carged gene comprising one or more chemical modifications and each strand of the double-stranded siNa comprises about 21 nucleotides. The siNa of the invention demonstrates antiarteriosclerotic, neuroprotective, or notropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian ctarget gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary diseases such as restences, atherescolerosis, Alzheimer's disease. Parkinson's disease, epilepsy, dementia, huntington's disease or amyotrophic lateral scalerosis. Purthermore, the sina may be utilised for gene therapy applications. The current sequence is that of the sina DNA-RNA may brid of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                          New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antiarteriosclerotic; neuroprotective; nootropic; antiparkinsonian; anticonvulsant; pulmonary disease; restenosis; atherosclerrosis; alabeimer's; Parkinson's; epidessy; dementia; huntington's; amyotrophic lateral sclerosis; gene therapy; ss; DNA-RNA hybrid; hTR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                 몫
                                                 Fosnaugh
                                                                                                                                                                                                                                                                                                                                                                                                                        tch 4.2%; Score 19; DB 1; Length 23; al Similarity 73.7%; Pred. No. 1.9e+02; 14; Conservative 5; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    double-stranded short interfering nucleic acid; siNA;
                                                 Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 23 BP; 4 A; 3 C; 7 G; 2 T; 5 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hTR-targeted sinA DNA-RNA hybrid - SEQ ID 604.
                                                                                                                                                                        Example 24; SEQ ID NO 606; 593pp; English.
                                               Chowrira B,
                                               J, Beigelman L, Chowr
, Usman N, Thompson J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             300 GAAGAGTTGGGCTCTGTCA 318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 GAAGAGUUGGGCUCUGUCA 20
                        (SIRN-) SIRNA THERAPEUTICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2002US-0358580P.
2002US-0363124P.
2002US-0386782P.
2002US-0406784P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADG30038 standard; RNA; 23 BP.
15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-FEB-2003; 2003WO-US005028
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-FEB-2004 (first entry)
                                                                                    WPI; 2003-731676/69.
                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unidentified
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                                                 Mcswiggen J.
Jamison S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADG30038;
                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 266
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of the invention demonstrates antiarteriosclerotic, neuroprotective, noticipally antiparkinsonian and anticionvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary diseases such as restenosis, atherosclerosis, Alzheimer's disease, Parkinson's disease, epilepsy, dementia, huntington's disease or mayotrophic lateral sclerosis. Furthermore, the siNA may be utilised for gene therapy applications. The current sequence is that of the siNA DNA-
                                                                                                                                                                                                                                                                                                                        New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                 Fosnaugh K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human telomerase RNA gene (hTR) promoter specific primer hlle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4.2%; Score 19; DB 1; Length 23;
53.2%; Pred. No. 1.9e+02;
ve 7; Mismatches 0; Indels
                                                                                                                                                             Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 23 BP; 4 A; 6 C; 2 G; 2 T; 7 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 24; SEQ ID NO 604; 593pp; English.
                                                                                                                                                             Chowrira B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RES CAMPAIGN TECHNOLOGY.
                                                                                                                                                                                               Thompson J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              146 TICCACCGIICATICIAGA 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ::||||||::||::||:
UUCCACCGUUCAUUCUAGA 20
                                                                                                 (SIRN-) SIRNA THERAPEUTICS INC.
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09-SEP-2002; 2002US-0409293P.
15-JAN-2003; 2003US-0440129P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAZ07305 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22-OCT-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity 63.2
nes 12; Conservative
                                                                                                                                                                    Beigelman
                                                                                                                                                                                               Usman N,
                                                                                                                                                                                                                                                            WPI; 2003-731676/69
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                                                                                                                                                                 Mcswiggen J,
Jamison S, t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-JAN-1998;
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US09436060A-16.rng.sl

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Disclosure, Fig 6; 109pp; English.
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                                                                                                         The invention relates to promoter regions from mouse and human telomerase

RNA (TR) component genes. The TR gene promoter can be linked to a

Reterologous gene, especially a gene encoding a cytoroxin, for therapy of

cancer, especially neoplasias. The telomerase is necessary for the

unrestricted proliferative capacity of many human cancers. Mutation or

dysregulation of the telomerase expression pathway may cause reactivation

or upregulation of telomerase expression pathway may cause reactivation

or upregulation of telomerase expression pathway may cause reactivation

or upregulation of telomerase expression pathway may cause

identified in the methods, can be used to block transcription from the TR

gene promoter through interaction of the 5' regulatory sequences. These

substances, e.g. antisense oligomotleotides, transcription factors,

peptide nucleic acids and factors that disrupt signal transduction, are

useful for cancer therapy. In particular, gene therapy vectors sequence

c useful for cancer therapy. In particular, gene therapy vectors and

c lespecially pGT62-codAupp) comprising the promoter and a viral thymidine

kinase gene can be controlled or treated. Direct down-regulation of

telomerase RNA gene through manipulation of transcription factors may be

effective anticancer therapy and the cloning of the hTR gene promoter

c allows the analysis of therapuetic molecules which modulate hTR promoter

activity. Sequences AAZ07656-221 regules

and mutagenisis of human TR gene (HTR) promoter region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                    Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; terc; gene therapy; thymidine kinase gene; anticancer therapy; mouse;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.2%; Score 18.8; DB 1; Length 22; 90.9%; Pred. No. 1.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 22 BP; 3 A; 4 C; 12 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mouse telomerase RNA gene specific primer mTRrl.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (CANC-) CANCER RES CAMPAIGN TECHNOLOGY.
                                                                                   Disclosure; Fig 12; 109pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 egeccredeadedeararedecc 22
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Les 20; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1999-479183/40.
WPI; 1999-479183/40
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                                                     gene therapy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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The invention relates to promoter regions from mouse and human telomerase RNA (TR) component genes. The TR gene promoter can be linked to a neterologous gene, especially a gene encoding a cytotoxin, for therapy of cancer, especially neoplasias. The telomerase is necessary for therapy of cancer, especially neoplasias. The telomerase is necessary for the concer, especially neoplasias. The telomerase is necessary for the concernation of the telomerase repression pathway may cause reactivation or upregulation of the telomerase expression in cancer. Substances.

CC dentified in the methods, can be used to block transcription from the TR dentified in the methods can be used to block transcription from the TR cancer through interaction of the 5' regulatory sequences. These substances, e.g. antisense oligonucleotides, transcription factors, peptide nucleic acids and factors that disrupt signal transduction, are useful for cancer therapy. In particular, gene therapy vectors

CC useful for cancer therapy in particular, gene therapy vectors

CC useful for cancer therapy. In particular, gene therapy vectors

CC useful for cancer therapy in particular, gene therapy vectors

CC especially pGF2-codatpp) comprising the promoter and a viral thymidine

CC telomerase RNA gene through manipulation of transcription factors may be effective anticancer therapy and the cloning of the hTR permoter

CC allows the analysis of therapeutic molecules which modulate hTR promoter

CC activity. Sequences AASOF68195 represent PCR primers for amplifying
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; terc; gene therapy; thymidine kinase gene; anticancer therapy; mouse;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 6 A; 5 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mouse telomerase RNA gene specific primer mTRrl.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CANC-) CANCER RES CAMPAIGN TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      activity. Sequences AAZU/b81-95 repress
mouse TR gene (terc) promoter sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20 TTCTCGCTGACTTCCAGCGG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-OCT-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 95.0
Matches 19; Conservative
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unrestricted proliferative capacity of many human cancers. Mutation or dysragulation of the telomerase repression pathway may cause reactivation of dysragulation of telomerase repression pathway may cause reactivation or upregulation of telomerase expression pathway may cause reactivation or upregulation of telomerase expression in cancer. Substances, edentified in the methods, can be used to block transcription from the TR gene promoter through interaction of the 5' regulatory sequences. These substances, e.g. antisense oligonucleotides, transcription factors, peptide nucleic acids and factors that disrupt signal transduction, are useful for cancer therapy. In particular, gene therapy vectors (especially prof62-codhupp) comprising the promoter and a viral thymidine kinase gene can be used to convert a prodrug, e.g. gancyclovir, so that neoplasia can be used to convert a prodrug, e.g. gancyclovir, so that coplasia can be used to convert a prodrug, e.g. gancyclovir, so that neoplasia can therapy and the cloning of transcription factors may be effective anticancer therapy and the cloning of the hTR gene promoter allows the analysis of therapeutic molecules which modulate hTR promoter activity. Sequences AAZ07681-95 represent PCR primers for amplifying
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Telomerase; human; cancer; diagnosis; melanoma; skin cancer; leukaemia; neuroblastoma; breast carciinoma; octoracciona; orecoma; osteosarcoma; smooth muscle cell hyperplasia; stem cell proliferation; Wilm's tumour; stem cell differentiation; organ fegeneration; organ differentiation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          products for
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4.1%; Score 18.4; DB 1; Length 20; 95.0%; Pred. No. 1.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gene (terc) promoter sequence
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97US-0053018P.
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97US-0054642P.
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telomerase of the invention. Primers that amplify the telomerase coding sequence can be used in a method for diagnosing cancer in a patient. The telomerase can be used for detection, diagnosis and drug screening. Inhibitors of telomerase activity can be used to treat cancers such as melanomas, other skin cancers, neuroblastomas, breast carcinomas, colon carcinomas, leukaemias, lymphomas, osteosarcomas or smooth muscle cell hyperplasias or skin growths. Enhancers of telomerase may be used to be activity and differentiation (expansion of haematopoietic stem cells could be administered in the bone marrow transplant context). As well, many tissues have stem cells. Proliferation of these cells may be useful in wound healing, hair growth, treatment of disease such as Wilm's tumour, organ regeneration or differentiation after injury or diseases, nerve cell or brain cell growth following
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New peptide nucleic acid (PNA) compounds that inhibit telomerase activity in mammalian cells is useful as probes to detect the RNA component of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note= "Peptide nucleic acid molecule, where N-(2-aminoethyl)glycine units are linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Peptide nucleic acid; PNA; telomerase; ribonucleoprotein enzyme; cance inhibitor; neoplasia; neurodegenerative disease; aging; hyperplasia; AIDS; HIV; fungal inflection; forensic identification; detect; tumour; paternicy testing; ss.
                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                             Score 18; DB 1; Length 18;
Pred. No. 1.7e+02;
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                                                                                                                                                                                                                                                               Sequence 18 BP; 1 A; 2 C; 12 G; 3 T; 0 U; 0 Other;
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100.0%; Pred. No. ...
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Best Local Similarity luv...
Best Local Similarity luv...
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controlled sequence contained within the RNA component of the enzyme. The nucleotide sequence contained within the RNA component of the enzyme. The invention relates to PNA molecules having a sequence of no more than 25 bases, which include the sequence of TAPAGG. The uncharged nature of the PNA cackbone increases the melting temperature of associating strands, increases the rate of association with targeted nucleic acids, and affords greater resistance of degradation by proteases or nucleases. The therapeutic PNAs may be used for treating disease conditions such as cancers, neoplasia, hyperplasia, neurodegenerative diseases, aging, human immunodeficiency virus (HIV) infection/AIDS (acquired immunodeficiency syndrome) and associated pathologies, fungal infections, and other cartivity, in combination with antineoplastic and other cytocated capants, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization cytostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization of individuals, e.g. paternity testing, based on hTR gene restriction of individuals, e.g. paternity testing, based on hTR gene restriction of individuals, e.g. paternity testing, based on hTR gene restriction of individuals, as probes to detect the RNA component of a mammalian telomerase and as inhibitors of telomerase activity. The method of the present invention allows are an early are an early and an early and an early and an early are the detected with increased conditions cander and as allows cancerous conditions to be detected with increases and as allows cancerous conditions to be detected with increase and as allows cancerous conditions to be detected with increase and as an early and an e
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//note= "This sequence is a peptide nucleic acid, i.e. it
contains a polyamide backbone instead of a deoxyribose
backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  possibly at an earlier stage, before cells are detected as cancerous based on pathological characteristics. The diagnostic and prognostic methods of the present invention can be used to detect an immortal or neoplastic cell or tumour tissue or cancer of any origin, provided the cell expresses telomerase activity and its RNA component
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PNA 27 inhibiting human and mammalian telomerase activity.
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Pred. No. 1.7e+02;
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Local Similarity 100.0%; Pr
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AAS15430 standard; DNA; 18 BP
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modified_base
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The present invention relates to peptide nucleic acids (PNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in mammalian cells by hybridising to the RNA component of mammalian cells by hybridising to the RNA component of mammalian cells by hybridising to the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polynucleotide having the human telomerase RNA component (hTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the hTR gene RFLP pattern. The CNA can be further used for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmacceuticals (such as antimeoplastic or cytostatic agents) can be used for treating neoplasia, hyperplasia, human immunodeficiency virus (HTV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomere metabolism or telomerase curviews.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                         New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; gene therapy; thymidine kinase gene; anticancer therapy; human; mutagenesis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human telomerase RNA gene (hTR) promoter specific primer h112b.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 18 BP; 2 A; 5 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                        Wright WE, Piatyszek MA, Corey DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (CANC-) CANCER RES CAMPAIGN TECHNOLOGY.
                                                                                                                                                                                                                                                           Claim 7; Col 73; 46pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      48 AACCCTAACTGAGAAGGG 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18 AACCCTAACTGAGAAGGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP
               97US-00838545.
96US-00630019.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22-OCT-1999 (first entry)
                                                 (TEXA ) UNIV TEXAS SYSTEM.
                                                                                                                        WPI; 2001-638024/73
09-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
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                 09-APR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         activity.
invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAZ07303;
                                                                                          Shay JW,
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Matches
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                                                                                                                                  The invention relates to promoter regions from mouse and human telomerase
RNA (TR) component genes. The TR gene promoter can be linked to a
cheterologous gene, sepecially a gene encoding a cytotoxin, for therapy of
cancer, especially nooplasias. The telomerase is necessary for the
unrestricted proliferative capacity of many human cancers. Mutation or
dysregulation of the telomerase repression pathway may cause reactivation
of telomerase expression pathway may cause reactivation
cor upregulation of telomerase expression in cancer. Substances,
identified in the methods, can be used to block transcription from the TR
gene promoter through interaction of the 5' regulatory sequences. These
cubstances, e.g. antiense oligonucleotides, transcription factors,
peptide nucleic acids and factors that disrupt signal transduction, are
useful for cancer therapy. In particular, gene therapy vectors
(especially pGT62-codAupp) comprising the promoter and a viral thymidine
kinase gene can be used to convert a prodrug, e.g. gancyclovir, so that
neoplasia can be used through manipulation of transcription factors may be
telomerase RNA gene through manipulation of transcription factors may be
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                                                             Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                                                                                                                                                                                                                                                effective anticancer therapy and the cloning of the hTR gene promoter allows the analysis of therapeutic molecules which modulate hTR promoter activity. Sequences AAZ07696-321 represent PCR primers used in cloning and mutagenisis of human TR gene (HTR) promoter region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RNA component; human telomerase; antisense oligonucleotide; infection; neuroblastoma; bladder cancer; colon cancer; prostate cancer; cancer; contraception; sterilisation; immunosuppression; therapeutic; hTR; immune system down-regulation; anti-inflammatory therapy; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RNA component of human telomerase (hTR) antisense oligo RP2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3.8%; Score 17.2; DB 1; Length 22;
86.4%; Pred. No. 2.5e+02;
ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pruzan R, Weinrich SL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 22 BP; 2 A; 4 C; 12 G; 4 T; 0 U; 0 Other;
                                                                                                           Disclosure; Fig 12; 109pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15 GGGCCTGGGAGGGGTGGTGGCC 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 égécérégéraAgérégéécé 22
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96US-00770565.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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Best Local Similarity 86.4
Matches 19; Conservative
                               WPI; 1999-479183/40.
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                                                                             gene therapy.
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20-DEC-1996;
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 Keith WN;
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Sequences shown in AAV41169 to AAV41181 represent antisense
oligonucleotides to the RNA component of human telomerase (hTR). These
antisense oligonuclectides specifically hybridise to a nucleotide
caquence within an accessible region of the hTR, but that does not
hybridise to a sequence within the template region of hTR. These
cligonucleotides may specifically be used for detection of an RNA
component of human telomerase in a sample. This is useful for diagnosing
cancer (especially neuroblastoma, bladder, colon and prostate cancer),
and providing prognosis for a cancer patient. The inhibitory
cligonucleotides can inhibit the telomerase activity level in a cell by
interfering with transcription of the RNA component, decreasing the half-
clife of the telomerase RNA component transcript, inhibiting assembly of
the RNA component into the telomerase notivery, no tinhibiting telomerase activity in both cultured cells and in
be used for inhibiting telomerase activity in both cultured cells and in
clels in vivo. They can be used in therapeutics for tracting or
munosuppression, and for selectively down-regulating specific branches
cof the immune system, e.g. a specific subset of T-cells, in anti-
inflammatory therapies or for treating infections by, e.g. yeast,
cancer for some and the seading infections by, e.g. yeast,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
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     - used for detecting
New polynucleotide(s) anti:sense to human telomerase - used for detecting or inhibiting human telomerase, e.g. for treating cancers, contraception, immuno-suppression or treating infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  acyl-coenzyme A synthetase 1; ACS1; diabetes; obesity; metabolic syndrome X; cardiovascular disorder; cancer; infection; inflammation; tumour; antisense; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3.8%; Score 17; DB 1; Length 17; 100.0%; Pred. No. 1.9e+02; ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 4 A; 6 C; 6 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                             Claim 11; Page 65; 80pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         177 TGTCAGCTGCTGGCCCG 193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP.
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Matches 17; Conservative
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ADK20555/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TARREST STATEMENT OF THE STATEMENT OF TH
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18-NOV-2004
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                                                                                           Query Match
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                                                                                                           Best Loca
Matches
                                                                                                                                                                                                                RESULT 277
886666
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                                                 The invention relates to an antisense compound targeted to a nucleic acid molecule encoding acyl-coenzyme A synthetase 1 (ACS1). The antisense compound specifically hybridises with and inhibits the expression of ACS1. The antisense oligonucleotides or compounds are useful for inhibiting the expression of acyl-coenzyme A synthetase 1 (ACS1), and for treating the expression of acyl-coenzyme A synthetase 1 (ACS1), and for treating diseases or conditions associated with aberrant expression of ACS1, e.g. diabetes, obestry, metabolic syndrome X, cardiovascular disorder or cancer. The antisense compounds are also useful as research reagents and kits, or in diagnostic, therapeutic and prophylactic applications, e.g. to prevent or delay infection, inflammation or tumour formation. The present sequence represents an acyl-coenzyme A synthetase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to an antisense compound targeted to a nucleic acid molecule encoding acyl-coenzyme A synthetase 1 (ACS1). The antisense compound specifically hybridises with and inhibits the expression of ACS1. The antisense oligonuclectides or compounds are useful for inhibiting the expression of acyl-coenzyme A synthetase 1 (ACS1), and for treating diseases or conditions associated with aberrant expression of ACS1, e.g. diabetes, obesity, metabolic syndrome X, cardiovascular disorder or cancer. The antisense compounds are also useful as research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New antisense compounds targeted to nucleic acid molecules encoding acyl-
coenzyme A synthetase 1 (ACS1), useful for treating diseases or
conditions associated with aberrant expression of ACS1, e.g. diabetes,
                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Acyl-coenzyme A synthetase 1, ACS1, antisense oligonucleotide #566.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 acyl-coenzyme A synthetase 1; ACS1; diabetes; obesity; metabolic syndrome X; cardiovascular disorder; cancer; infection; inflammation; tumour; antisense; ss.
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                                                                                                                                                                                                                                                     Score 16.4; DB 1; Length 20;
Pred. No. 2.6e+02;
0; Mismatches 1; Indels
                                                                                                                                                                                                                              Sequence 20 BP; 1 A; 12 C; 4 G; 3 T; 0 U; 0 Other;
                          SEQ ID NO 632; 940pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 3; SEQ ID NO 566; 940pp; English.
                                                                                                                                                                                                   1, ACS1, antisense oligonucleotide
                                                                                                                                                                                                                                                                                                             349
                                                                                                                                                                                                                                                                                                                                                                                                          BP
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                                                                                                                                                                                                                                                         3.6%;
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                                                                                                                                                                                                                                                                                                              CGGGGGCGAGGCGAGGT
                                                                                                                                                                                                                                                                                                                                        CGAGGCGAGGCGAGGT
                                                                                                                                                                                                                                                                                                                                                                                                          ADK20489 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                  17; Conservative
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                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    obesity or cancer.
 obesity or cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                              332
                                                                                                                                                                                                                                                                                                                                        13
                                                                                                                                                                                                                                                                                                                                                                                                                                     ADK20489;
                                                                                                                                                                                                                                                          Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New antisense compounds targeted to nucleic acid molecules encoding acylocarzyme A synthetase 1 (ACS1), useful for treating diseases or conditions associated with aberrant expression of ACS1, e.g. diabetes,
reagents and kits, or in diagnostic, therapeutic and prophylactic applications, e.g. to prevent or delay infection, inflammation or tumour formation. The present sequence represents an acyl-coenzyme A synthetase 1, ACS1, antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Acyl-coenzyme A synthetase 1, ACS1, antisense oligonucleotide #713.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     acyl-coenzyme A synthetase 1; ACS1; diabetes; obesity;
metabolic syndrome X; cardiovascular disorder; cancer; infection;
inflammation; tumour; antisense; ss.
                                                                                                                                                                                                                                                                                                                      .
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0
                                                                                                                                                                                                                                                 3.6%; Score 16.4; DB 1; Length 20;
94.4%; Pred. No. 2.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3.6%; Score 16.4; DB 1; Length 20;
14.4%; Pred. No. 2.6e+02;
                                                                                                                                                                                                                                                                                                                      1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
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                                                                                                                                                                                Sequence 20 BP; 1 A; 12 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                      0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1, ACS1, antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            332 CGGGGGGGGGGGGGT 349
                                                                                                                                                                                                                                                                                                                                                                                              332 CGGGGGCGAGGCGAGGT 349
                                                                                                                                                                                                                                                                                                                                                                                                                                                           18 ccacccacccacccacct 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-AUG-2003; 2003WO-US025389.
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                                                                                                                                                                                                                                                                                     ilarity 94.4%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 3.6%;
Best Local Similarity 94.4%;
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADK20636 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-203782/19.
                                                                                                                                                                                                                                                                                     Local Similarity
hes 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               obesity or cancer.
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The inventors describe the production of mammalian cells expressing prohormone convertase which facilitates the processing of prohormone procursors to active hormones. More specifically the cleavage site is the prohormone convertase cleavage site given in ARR37632-35. In order to define murine prohormone convertase 1 specifically prorelaxin alanine mutations of the basic residues K and R were constructed. The following mutations were tested for substrate specificity to murine prohomone convertase: 1.4, 2.4, 3.2, 4.3, 7.2 and 8.6 (see AAQ43256-63; AAR37619-26). (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence encoding the C-chain/A-chain junction prorelaxin (Rx) mutant Rx 2.4 at the prohormone convertage cleavage site.
                                                                                              Sequence encoding the C-chain/A-chain junction prorelaxin (Rx) mutant Rx 1.4 at the prohormone convertase cleavage site.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Heterologous polypeptide factor prepn. - by introducing into polypeptide factor dependent host cell nucleic acid, and then culturing host cell,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Prohormone convertage; enzyme; cleavage; prorelaxin; hormone;
                                                                                                                                                       Prohormone convertage; enzyme; cleavage; prorelaxin; hormone;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    match 3.6%; Score 16.2; DB 1; Length 21; Local Similarity 85.7%; Pred. No. 2.8e+02; ne 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 10 A; 4 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Marriott D;
                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      156 CATTCTAGAGCAAACAAAAA 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 CATTCTAGAGCAAAGAGACAA 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example; Page 54; 134pp; English.
                                                                                                                                                                                                                                                                                                                                                                                          92WO-US010621.
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92US-00887265
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAQ43258 standard; DNA; 21
                                      (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1993-197065/24.
P-PSDB; AAR37620.
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24-SEP-1993
                                      25-MAR-2003
24-SEP-1993
                                                                                                                                                                                                                                                                                                                                                     10-JUN-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gorman CM,
                                                                                                                                                                                                Synthetic.
AAQ43257;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2 Primers (see AAV27886 and AAV27887) are used in the PCR amplification of a 520 bp fragment of telomerase genomic DNA from HeLa cells. The PCR product was used to prepare 2 DNA constructs, each containing the T7 promoter. One construct (for primers see AV27888-89) contained DNA which would generate sense strand human telomerase RNA in a transcription reaction. The other construct (for primers see AAV27890-91) contained DNA which would generate antisense strand human telomerase RNA in a transcription reaction. The human telomerase RNA in a transcription reaction. The human telomerase RNA was used for in vitro assays to demonstrate that wild-type TP2 and sense telomerase RNA are required for telomerase activity, Novel TP2 can be used to develop products that regulate telomerase activity, useful e.g. for treating fancer, AIDS and ageing disorders. (Updated on 25-MAR-2003 to correct PR field.)
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                                                                                                                                                                                                                                                                                              PP2; human; telomerase protein 2; cancer; AIDS; ageing; therapy; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New nucleic acid encoding human telomerase protein-2 - used for regulating telomerase activity, e.g. for treating cancer or acquired
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 1 A; 5 C; 13 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 8; Page 90; 150pp; English.
                                                                                                                                                                                                                                                         Human telomerase gene PCR primer.
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                                                                                                                    AAV27886 standard; DNA; 21 BP
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20 CGAGGGCGAGGCGAGGT
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                                                                                                                                                                                                (revised)
(first entry)
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(AMGE-) AMGEN CANADA INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1998-297946/26.
                                                                                                                                                                                                                                                                                                                                                     Synthetic.
Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-JUN-1997;
16-OCT-1997;
                                                                                                                                                                                                                                                                                                                                                                                                              WO9821343-A1
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                                                                                                                                                                                              25-MAR-2003
12-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-NOV-1996;
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                                                                                                                                                                                                                                                                                                                  primer; ss
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AAV27886

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RESULT 279 AAQ43257 ID AAQ432 XX

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(GETH ) GENENTECH INC
                                       WPI; 1994-303031/37.
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                                                  P-PSDB; AAR60585
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
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04-AUG-1997;
09-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9901560-A1
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21-JUL-1997;
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                      Gorman CM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                      AAX18326;
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                                                                                                                                                                                                                                                        The inventors describe the production of mammalian cells expressing prohormone convertase which facilitates the processing of prohormone products to active hormones. More specifically the cleavage site is the prohormone convertase cleavage site given in ARR3762-35. In order to define murine prohormone convertase 1 specifically prorelaxin alanine mutations of the basic residues K and R were constructed. The following mutations were tested for substrate specificity to murine prohomone convertase: 1.4, 2.4, 3.2, 4.3, 7.2 and 8.6 (see AAQ43256-63; AAR37619-26). (Updated on 25-WAR-2003 to correct PN field.)
                                                                                                                                                                                                     Heterologous polypeptide factor prepn. - by introducing into polypeptide factor dependent host cell nucleic acid, and then culturing host cell,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           prohormone; convertase; insulin; proinsulin; factor; growth; precursors;
transgenic; mammal; prorelaxin; cleavage; ss.
                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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0
                                                                                                                                                                                                                                                                                                                                                                       3.6%; Score 16.2; DB 1; Length 21;
85.7%; Pred. No. 2.8e+02;
iive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rx 2.4 prorelaxin C/A chain junction mutant cleavage site.
                                                                                                                                                                                                                                                                                                                                                    Sequence 21 BP; 10 A; 4 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                        Gorman CM, Groskreutz DJ, Marriott D;
                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                          CATTCTAGAGCAAACAAAAA 176
                                                                                                                                                                                                                                                                                                                                                                                                                            CATTCTAGAGCAAAGAGACAA 21
                                                                                                                                                                                                                                           Example; Page 54; 134pp; English.
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                                                                                           92WO-US010621
                                                                                                             91US-00803631
                                                                                                                       92US-00887265
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(first entry)
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                                    /*tag=
                                                                                                                                       (GETH ) GENENTECH INC.
                                                                                                                                                                           WPI; 1993-197065/24.
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                                                                                                                                                                                    P-PSDB; AAR37621
                                                       WO9311247-A1
                                                                                          04-DEC-1992;
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                                                                        10-JUN-1993
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26-APR-1995
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Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                           156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ71457;
                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 281
                                                                                                                                                                                                                                                                                                                                                                                       Matches
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A series of natural or mutated cDNAs for the prohormone convertase (PC) cleavage sites in human proralaxin protein between the C and A chains (AAQ71455-59 corresponding to AR860583-7) or in the B and C chains (AAQ71460-63 corresponding to AR860588-90). The natural cleavage sites contain two dibasic residues, Lys-Arg, both of which are required for cleavage. The mutant sites have replacements of either residues with an Ala. This confers specificity to murine PCI. In this particular mutant, the Lys residue at the fourth position of the cleavage site has been responsible for remodelling the reproductive tract prior to parturition. The cells containing both the prohormone convertase gene and the required precursor to be expressed can be injected into a mammal. (Updated on 25-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ò
                                                                                                                                                                                            Treating insulin-dependent disorders in mammals - by introducing a nucleic acid encoding a variant proinsulin into a host cell with a constitutive pathway of protein secretion, or a plasmid, and introducing the cell or plasmid to the mammal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          neuroblastoma; breast carcinoma; colon carcinoma; lymphoma; osteosarcoma; smooth muscle cell hyperplasia; stem cell proliferation; Wilm's tumour; stem cell differentiation; organ regeneration; organ differentiation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Telomerase; human; cancer; diagnosis; melanoma; skin cancer; leukaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 3.6%; Score 16.2; DB 1; Length 21; Best Local Similarity 85.7%; Pred. No. 2.8e+02; Matches 18; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 10 A; 4 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCR primer for telomerase coding sequence.
                                                                                                                                                                                                                                                                                                                                                                                             Example 3; Page 52; 117pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         156 CATTCTAGAGCAAACAAAAA 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 CATTCTAGAGCAAAGAGACAA 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP
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97US-0053018P.
97US-0053329P.
97US-0054642P.
97US-0058287P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98WO-US013835.
Groskreutz DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAX18326 standard; DNA; 21
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82333333
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                                                                                                                                  This sequence is a PCR primer for DNA encoding a truncated human telomerase of the invention. Primers that amplify the telomerase coding sequence can be used in a method for diagnosing cancer in a patient. The telomerase can be used for detection, diagnosis and drug screening. Inhibitors of telomerase activity can be used to treat cancers such as meanomas, other skin cancers, neuroblastoomas, breast carcinomas, colon carcinomas, leukaemias, lymphomas, osteosarcomas or smooth muscle cell hyperplasias or skin growths. Enhancers of telomerase may be used to be stimulate stem cells could be administered in the bone marrow transplant context). As well, many tissues have stem cells. Proliferation of these cells may be useful in wound healing, hair growth, treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                            New isolated vertebrate telomerase genes - used to develop products for treating cancers or for organ regeneration, nerve cell or brain cell growth following injury or bone marrow transplantation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                              disease such as Wilm's tumour, organ regeneration or differentiation after injury or diseases, nerve cell or brain cell growth following
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Peptide nucleic acid; PNA; cancer; telomerase; probe; hybridisation;
inhibitor; human telomerase RNA; hTR; PCR; oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New peptide nucleic acids hybridising to mammalian telomerase RNA -
to inhibit telomerase, for treating tumours and other proliferative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA oligonucleotide 3, used in the measurement of Tm values.
                                                                                                                                                                                                                                                                                                                                                                                                                         3.6%; Score 16.2; DB 1; Length 21; 85.7%; Pred. No. 2.8e+02; ive 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21 BP; 3 A; 5 C; 9 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Corey D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Piatyszek MA,
                                                                                                          Example 1, Page 43; 134pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         431 CAGGACTCGGCTCACACATGC 451
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 2; Page 49; 76pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CAGGACTCGGCTCACCACTGC 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 diseases, also for diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAT89247 standard; DNA; 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       09-APR-1997;
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                                                                                                                                                                                                                                                                                                                                                                  injury
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This is an oligonucleotide used in the measurement of Tm values and their complimentary peptide nucleic acids (PNAs), (e.g. AAT89236-T89239). PNAs

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New peptide nucleic acid (PNA) compounds that inhibit telomerase activity in mammalian cells is useful as probes to detect the RNA component of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Peptide nucleic acid; PNA; telomerase; ribonucleoprotein enzyme; cancer; inhibitor; neoplasia; neurodegenerative disease; aging; hyperplasia; AIDS; HIV; fungal infection; forensic identification; detect; tumour; paternity testing; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note= "Peptide nucleic acid molecule, where N-(2-aminoethyl)glycine units are linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
hybridise specifically to an RNA component of mammalian telomerase, and include the sequence GGG for specific hybridisation to the template region of this component. PNAs can be used as probes to detect the RNA component of mammalian telomerase and as inhibitors of telomerase activity, especially in the treatment of cancer
                                                                                                                                                                                              Gaps
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                                                                                                                                                  DB 1; Length 16;
2.1e+02;
                                                                                                                                                                                              0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Corey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PNA sequence #27 used to inhibit telomerase activity.
                                                                                                                    Sequence 16 BP; 5 A; 2 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Norton JC,
                                                                                                                                                        3.5%; Score 16; DB 100.0%; Pred. No. 2.1 active 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Shay JW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                      BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            97US-00838545.
                                                                                                                                                                                                                                                              TAACTGAGAAGGGCGT 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  96US-00630019
                                                                                                                                                                                                                                    68
                                                                                                                                                                                                                                                                                                                                                                      AAA37569 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Piatyszek MA,
                                                                                                                                                                                                                                    53 TAACTGAGAAGGGCGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (TEXA ) UNIV TEXAS SYSTEM
                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mammalian telomerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2000-292432/25.
                                                                                                                                                        Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  misc feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                            AAA37569;
                                                                                                                                                                                                                                                                                                                                  RESULT 284
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syndrome) and associated by abnormal telomere metabolism and other diseases characterized by abnormal telomere metabolism or telomerase cativity, in combination with antineoplastic and other cytocoxic or cytostatic agents, antifungal agents, and other cytocoxic or cytostatic agents, antifungal agents, and other nucleotides. PNAs may be compared for molecular diagnostics, labelled PNAs are used as hybridization probes to detect or quantitate polymucleotides having a human telomerase RNA (hTR) sequence. PNA probes are also used for forensic identification of individuals, e.g. paternity testing, based on hTR gene restriction fragment length polymorphism (RFLP) pattern. PNAs are also useful as probes to detect the RNA component of a mammalian telomerase and as inhibitors of telomerase activity. The method of the present invention possibly at an earlier stage, before cells are detected as cancerous conditions to be detected as cancerous based on pathological characteristics. The diagnostic and prognostic methods of the present invention can be used to detect an immortal or neoplastic cell or tumour tissue or cancer of any origin, provided the respresses telomerase activity and its RNA component. immunodeficiency virus (HIV) infection/AIDS (acquired immunodeficiency

Sequence 16 BP; 5 A; 2 C; 6 G; 3 T; 0 U; 0 Other;

.; 0 Gaps .; 0 3.5%; Score 16; DB 1; Length 16; 100.0%; Pred. No. 2.1e+02; cive 0; Mismatches 0; Indels 68 53 TAACTGAGAAGGGCGT Local Similarity 100. 16; Conservative Query Match Best Loca Matches ð

1 TAACTGAGAAGGGCGT 16 셤

AAS15447 standard; DNA; 16 BP RESULT 285 AAS15447

AAS15447;

Oligonucleotide #3 used in melting temperature studies of PNAs. (first entry) 14-FEB-2002

Mammalian; paternity testing; human telomerase RNA component; hTR gene RFLP pattern; cancer; inflammation; forensic; lymphoproliferative disease; autoimmune disease; hyperplasia; neurodegenerative disease; neoplasia; HIV; AIDS; cytostatic; human immunodeficiency virus; acquired immunodeficiency syndrome; telomere metabolism; anti-inflammatory; immunosuppressive; ss.

sapiens Synthetic. Ношо

US6294650-B1

25-SEP-2001,

99US-00349532. 08-JUL-1999;

96US-00630019. 97US-00838545. 09-APR-1996; 09-APR-1997;

(TEXA) UNIV TEXAS SYSTEM,

Shay JW, Wright WE,

Norton JC;

Piatyszek MA, Corey DR,

WPI; 2001-638024/73,

New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases.

Example 2; Col 34; 46pp; English.

The present invention relates to peptide nucleic acids (PNAs), comprising

mammalian cells by hybridising to the RNA component of mammalian cells by hybridising to the RNA component of mammalian cells by hybridising to the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to mammalian telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polynucleotide having the human telomerase RNA component (hrr) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the HTR gene RFLP pattern. The PNA can be further used for treating or identification of criminal suspects or unknown descendants based on the PTR gene RFLP pattern. The PNA can be further used for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases, neombination with other pharmaceuticals (such as antineoplastic or cytostatic againts) can be used for treating neoplasia, hyperplasia, human immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomere metabolism or telomerase activity. The present sequence representing a DNA oligonucleotide is complementary to some of the PNAs of the present invention, and is used ö The invention relates to promoter regions from mouse and human telomerase RNA (TR) component genes. The TR gene promoter can be linked to a heterologous gene, especially a gene encoding a cytotoxin, for therapy of cancer, especially neoplasias. The telomerase is necessary for the unrestricted proliferative capacity of many human cancers. Mutation or dysregulation of the telomerase repression pathway may cause reactivation or upregulation of telomerase expression in cancer. Substances, and human telomerase RNA gene promoters, useful for tumor specific Gaps sequence of 6-25 nucleobases, that inhibit telomerase activity in Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; gene therapy; thymidine kinase gene; anticancer therapy; human; ; 0 3.5%; Score 16; DB 1; Length 16; 100.0%; Pred. No. 2.1e+02; Live 0; Mismatches 0; Indels Human telomerase RNA gene (hTR) specific primer hTRg. Sequence 16 BP; 5 A; 2 C; 6 G; 3 T; 0 U; 0 Other; (CANC-) CANCER RES CAMPAIGN TECHNOLOGY, Disclosure; Fig 6; 109pp; English. in melting temperature studies BP. 1 TAACTGAGAAGGGCGT 16 53 TAACTGAGAAGGGCGT 68 98GB-00001902. 99WO-GB000308. AAZ07277 standard; DNA; 20 (first entry) Query Match 3.5 Best Local Similarity 100. Matches 16; Conservative WPI; 1999-479183/40. Homo sapiens. 22-OCT-1999 WO9938964-A2 29-JAN-1999; 29-JAN-1998; gene therapy PCR primer; 05-AUG-1999. AAZ07277; Synthetic Keith WN; RESULT 286 Mouse 셤 à

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identified in the methods, can be used to block transcription from the TR gene promoter through interaction of the 5' regulatory sequences. These bubstances, e.g. antisense oligonucleotides, transcription factors, peptide nucleic acids and factors that disrupt signal transduction, are useful for cancer therapy. In particular, gene therapy vectors (especially pGTG-codAupp) comprising the promoter and a viral thymidine kinase gene can be used to convert a prodrug, e.g. gancyclovir, so that neoplasia can be controlled or treated. Direct down regulation of telomerase RNA gene through manipulation of transcription factors may be effective anticancer therapy and the cloning of the hTR gene promoter allows the analysis of therapeutic molecules which modulate hTR promoter activity. Sequences AAZ07621-80 represents PCR primers for amplifying human TR gene (HTR) promoter sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           function of nucleic acid molecules encoding hypothetical tumour endothelial marker. The antisense compound can also be used as research tools and diagnostics. It can also be used as tools in differential and/or combinatorial analyses to elucidate expression patterns of a portion or the entire complement of genes expressed within cells and tissues. The compound can also be used for treating diseases or conditions associated with hypothetical tumour endothelial marker, preferably hyperproliferative disorder, e.g. colon cancer. The compound can also be used as prophylaxis, e.g. to prevent or delay infection,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New antisense compound targeted to a nucleic acid molecule encoding hypothetical tumor endothelial marker, useful for modulating expression of hypothetical tumor endothelial marker or for treating colon cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a compound targeted to a nucleic acid molecule encoding hypothetical tumour endothelial marker. The compound, particularly the antisense oligonucleotide is useful in modulating the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human tumour endothelial marker antisense oligonucleotide ISIS 208420.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  human; 88; antisense; hypothetical tumour endothelial marker; tumour; hyperproliferative disorder; colon cancer; infection; inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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0
                                                                                                                                                                                                                                                                                                                                                       3.5%; Score 16; DB 1; Length 20; 100.0%; Pred. No. 2.8e+02; ive 0; Mismatches 0; Indels
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Best Local Similarity 100.'
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-061309/06.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADH56499/
      8888888888888888888888888
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The invention relates to a compound targeted to a nucleic acid molecule encoding hypothetical tumour endothelial marker. The compound, particularly the antisense oligonucleotide is useful in modulating the function of nucleic acid molecules encoding hypothetical tumour endothelial marker. The antisense compound can also be used as research tools and diagnostics. It can also be used as tools in differential and/or combinatorial analyses to elucidate expression patterns of a portion or the entire complement of genes expressed within cells and tissues. The compound can also be used for treating diseases or conditions associated with hypothetical tumour endothelial marker, preferably hyperproliferative disorder, e.g. colon cancer. The compound can also be used as prophylaxis, e.g. to prevent or delay infection, inflammation or tumour formation. The present sequence represents a human human human endothelial marker target region.
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inflammation or tumour formation. The present sequence represents a human hypothetical tumour endothelial marker target region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New antisense compound targeted to a nucleic acid molecule encoding hypothetical tumor endothelial marker, useful for modulating expression of hypothetical tumor endothelial marker or for treating colon cancer.
                                                                                                                                                                                                                                                                                                                                                                                                           Human hypothetical tumour endothelial marker target region ISIS 126034.
                                                                                                                               Сарв
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Сарв
                                                                                                                                                                                                                                                                                                                                                                                                                                               human, 88; hypothetical tumour endothelial marker; tumour;
hyperproliferative disorder; colon cancer; infection; inflammation.
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                                                                                        3.5%; Score 15.8; DB 1; Length 20; 89.5%; Pred. No. 2.9e+02;
                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                    Sequence 20 BP; 1 A; 7 C; 4 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 8 A; 4 C; 7 G; 1 T; 0 U; 0 Other;
                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 15; SEQ ID NO 138; 102pp; English.
                                                                                                                                                                362 GGCCGCAGGAAGGAACG 380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  362 GGCCGCAGGAAGAGAACG 380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13
                                                                                                                                                                                         20 GGCCACAGGAAAGGAACG 2
                                                                                                                                                                                                                                                                                                 ВЪ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17-JUN-2002; 2002US-00174020
                                                                                                                                                                                                                                                                                               ADH56566 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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Best Local Similarity 89.5
Matches 17; Conservative
                                                                                                          Local Similarity 89.5
18 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Monia BP, Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-061309/06.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                        25-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18-DEC-2003.
                                                                                                                                                                                                                                                                                                                                  ADH56566;
                                                                                          Query Match
                                                                                                                             Matches
                                                                                                                                                                                                                                                          RESULT 288
                                                                                                                                                                                                                                                                               ADH56566
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AAT89233;

ADP27647;

RESULT 289

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This sequence is a novel peptide nucleic acid (PNA), which acts as an inhibitor of mammalian, preferably human, telomerase. The PNAs hybridise specifically to an RNA component of mammalian telomerase, and include the sequence GGG for specific hybridisation to the template region of this component. PNAs can be used as probes to detect the RNA component of mammalian telomerase and as inhibitors of telomerase activity, especially
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RBP-7; retinoblastoma binding protein-7; abnormal cell proliferation; diagnosis; therapy; cell differentiation; thyroid hyperplasia; psoriasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New peptide nucleic acids hybridising to mammalian telomerase RNA - used to inhibit telomerase, for treating tumours and other proliferative
                                                                                                                                                                                                                                                                             1. .17
/*tag= a
/#ote= "Sugar-phosphate backbone has been replaced by a
peptide backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Revised record issued on 21-OCT-2004 : Correction to feature table key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCR primer for human retinoblastoma binding protein-7 coding sequence.
                                                                                                                                                                 Peptide nucleic acid; PNA; cancer; telomerase; probe; hybridisation;
                                                                                                                       Peptide nucleic acid 9, targeted to mammalian telomerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 2 A; 5 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 15.4; DB 1;
Pred. No. 2.5e+02;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Piatyszek MA,
                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 9; Page 59; 76pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            diseases, also for diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                97WO-US005931.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             96US-00630019
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              48 AACCCTAACTGAGAAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17 AACCCTAACTGAGAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAZ87074 standard; DNA; 20
                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               in the treatment of cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-MAY-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity 94.1
nes 16; Conservative
                                                            (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Wright WE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1997-512647/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GERO-) GERON CORP
                                                                                                                                                                                          .nhibitor; ss.
                                                                                                                                                                                                                                                                               Key
modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                             WO9738013-A1
                                                            21-OCT-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16-0CT-1997.
                                                                                12-MAY-1998
                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAZ87074;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Shay JW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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à
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This invention relates to a novel method concerning prognostic markers associated with EGFR (epidermal growth factor receptor) positive cancer. Specifically, it refers to a gene expression profiling method that can provide a prediction as to whether a patient is likely to respond well to treatment with an EGFR inhibitor. The present invention describes the current cone gene selected from the group of CD4403, CD4406, DR5, GRO1, KRP17, LAWC2 or their products thereof. It further provides a cDNA microarray containing named genes that represent prognostic transcripts which are useful for determining whether a patient diagnosed with an EGFR capressing head or neck cancer or colon cancer exhibits elevated or decreased expression levels of these genes compared to normal. As such, these methods are also useful for prognosing or predicting the likelihood of cancer-attributable death or prognosing or predicting the likelihood or cancer-attributable death nor progression, including recurrence and metastatic spread of a neoplastic disease, as well as drug resistance. This polynucleotide sequence is a human PCR amplicon DNA sequence used as prognostic cancer marker, given in an exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Prognosing a patient with EGFR-expressing colon cancer comprises subjecting a sample comprising EGFR-expressing cancer cells to quantitative analysis of the expression level of the RNA transcript of at least one gene e.g., CD44v3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Gaps
                                                                                                                                                                                                                                              TERC; human; PCR amplicon; ds; prognostic marker; EGFR; epidermal growth factor receptor; cancer; gene expression profiling; microarray; head and neck cancer; colon cancer; metastatic spread; neoplastic disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 15.8; DB 1; Length 79;
Pred. No. 2.5e+02;
0; Mismatches 27; Indels
                                                                                                                                                                                                                   Human TERC DNA used as a cancer prognostic marker SeqID 84.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 79 BP; 16 A; 26 C; 27 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Baselga J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 55; SEQ ID NO 84; 113pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GENO-) GENOMIC HEALTH INC.
(VALL-) VALL HEBRON UNIV HOSPITAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Shak S,
                                                                                   ADP27647 standard; DNA; 79 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-NOV-2003; 2003WO-US036777.
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Local Similarity 54.2%;
Les 32; Conservative
                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cronin MT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-420643/39.
                                                                                                                                                                                                                                                                                                                                                                                                               WO2004046386-A1.
                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                          26-AUG-2004
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Norton JC;

Corey D,

ó;

Gaps

0;

Indels

AAT89233/c ID AAT89233 standard; DNA; 17 BP.

RESULT 290

65

Query Match

Best Loc Matches

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Length 17;

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14-AUG-2003; 2003WO-US025389.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADK20828;
                                                                                                                                                                                               ROSB SA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ROBB SA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADK20828,
   THE CONTRACT OF THE CONTRACT O
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ð
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This sequence represents a PCR for the retinoblastoma binding protein-7 (RBP-7) genomic sequences are useful for the RBP-7 coding sequence and regulatory sequences are useful for the recombinant production of the protein and for expressing heterologous nucleic acids. Primers and probes derived from the RBP-7 nucleotide sequence (such as this sequence) are useful for DNA amplification and detection methods. RBP-7 biallelic markers (see AZ86993-Z87034) are useful for diagnosis of disease related to alteration in the regulation or in the coding regions of the RBP-7 gene and for prognosis/diagnosis of an eventual treatment with therapeutic agents, especially agents acting on pathologies involving abnormal cell proliferation and/or differentiation, these include thyroid hyperplasia, psoriasis, benign prostate hypertrophy, cancers, including breast cancer, sercomas and other neoplasms, bladder cancer, colon cancer, lung cancer, prostate cancer, various leukaemias,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     novel nucleic acid and polymorphic markers used for diagnosis of diseases, especially those involving abnormal cell proliferation and differentiation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Acyl-coenzyme A synthetase 1, ACS1, antisense oligonucleotide #727.
benign prostate hypertrophy; cancer; sarcoma; neoplasm; leukaemia;
lymphoma; biallelic marker; PCR primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         acyl-coenzyme A synthetase 1; ACS1; diabetes; obesity;
metabolic syndrome X; cardiovascular disorder; cancer; infection;
inflammation; tumour; antisense; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              and lymphomas. RBP-7 antibodies are useful as diagnostic agents
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 9 A; 3 C; 4 G; 4 T; 0 U; 0 Other,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 16; Page 206; 223pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 CAAACAATAAATGTCAG 17
                                                                                                                                                                                                                                                                                                                                                      98US-0091315P.
                                                                                                                                                                                                                                                                                          99WO-IB001242
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-117170/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
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                                                                                                                                                             WO200000607-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                       (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bougueleret L;
                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                          30-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                      30-JUN-1998;
10-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18-NOV-2004
                                                                                                                                                                                                                             06-JAN-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26-FEB-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADK20650/c
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                                                                                                                                                                                                                                                                                                             New antisense compounds targeted to nucleic acid molecules encoding acyl-coenzyme A synthetase 1 (ACS1), useful for treating diseases or conditions associated with aberrant expression of ACS1, e.g. diabetes,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New antisense compounds targeted to nucleic acid molecules encoding acyl-coenzyme A synthetase 1 (ACS1), useful for treating diseases or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    acyl-coenzyme A synthetase 1; ACS1; diabetes; obesity; metabolic syndrome X; cardiovascular disorder; cancer; infection; inflammation; tumour; antisense; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 2 A; 13 C; 2 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 3; SEQ ID NO 727; 940pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              349
14-AUG-2002; 2002US-0403591P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-AUG-2003; 2003WO-US025389
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 94.1%;
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              333 GGGGGGGAGGCGAGGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20 GAGGGGGGGGGGGT
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                                                                             (PHAA ) PHARMACIA CORP
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                                                                                                                                                                                                                                        WPI; 2004-203782/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                       obesity or cancer
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Wed Aug

Length 20;

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3.4%; Score 15.2; DB 1;
85.0%; Pred. No. 3.2e+02;
iive 0; Mismatches 3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1, ACS1, antisense oligonucleotide.
                                                     313 CTGTCAGCCGCGGGTCTCTC 332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      328 CTCTCGGGGCGAGGCGAG 347
                                                                                1 CTGTCTGCCACGGGTTTCTC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP.
                                                                                                                                                                                                                                                                                                                                                                                                                  14-AUG-2002; 2002US-0403591P.
                                                                                                                                                                                                                                                                                                                                                                                       14-AUG-2003; 2003WO-US025389.
                                                                                                                                                 ADK21153 standard; DNA; 20
                                                                                                                                                                                                     (first entry)
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               Local Similarity 85.0
nes 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2004-203782/19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 obesity or cancer.
                                                                                                                                                                                                                                                                                                                                    WO2004016749-A2.
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                                                                                                                                                                                                                                                                                                           Synthetic
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    Query Match
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                 Best Loca
Matches
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Matches
                                                                                                                        RESULT 295
                                                                                                                                   ADK21153/
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                                                                The invention relates to an antisense compound targeted to a nucleic acid molecule encoding acyl-coenzyme A synthetase 1 (ACS1). The antisense compound specifically hybridises with and inhibits the expression of ACS1. The antisense oligonuclectides or compounds are useful for inhibiting the expression of acyl-coenzyme A synthetase 1 (ACS1), and for treating diseases or conditions associated with aberrant expression of ACS1, e.g. diabetes, obesity, metabolic syndrome X, cardiovascular disorder or cancer. The antisense compounds are also useful as research reagents and kits, or in diagnostic, therapeutic and prophylactic applications, e.g. to prevent or delay infection, inflammation or tumour formation. The present sequence represents an acyl-coenzyme A synthetase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel rat pheochromocytoma PC12 cell line useful as cellular model of Huntington's disease, comprising an expression cassette containing a DNA encoding a truncated mutant huntingtin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention provides a rat or mouse pheochromocytoma cell line stably transfected with an expression cassette encoding a truncated mutant buntingtin. This can be used in the study of the pathological mechanism in Huntington's disease and enables the development of therapies and diagnostic techniques for the disease
    conditions associated with aberrant expression of ACS1, e.g. diabetes,
                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; huntingtin; Huntington's disease; immunogen; fusion protein;
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0
                                                                                                                                                                                                                                                                  3.4%; Score 15.4; DB 1; Length 20;
4.1%; Pred. No. 3.1e+02;
ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rat/human glutamate transporter GLAST PCR primer #2.
                                                                                                                                                                                                                                         Sequence 20 BP; 1 A; 12 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 1 A; 7 C; 5 G; 7 T; 0 U; 0 Other;
                                         Claim 3; SEQ ID NO 905; 940pp; English.
                                                                                                                                                                                                                   1, ACS1, antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Page 21; 55pp; English.
                                                                                                                                                                                                                                                                                                                         332 CGGGGGCGAGGCGAGG 348
                                                                                                                                                                                                                                                                                                                                                                                                                    BP
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                                                                                                                                                                                                                                                                                 94.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                               16; Conservative
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Best Local Similarity
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                   obesity or cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                             AAC67060;
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                                                                                                                                                                                                                                                                                                Matches
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Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Acyl-coenzyme A synthetase 1, ACS1, antisense oligonucleotide #1230.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            acyl-coenzyme A synthetase 1; ACS1; diabetes; obesity;
metabolic syndrome X; cardiovascular disorder; cancer; infection;
inflammation; tumour; antisense; ss.
0;
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Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 296
AAT89229/c
ID AAT89229 standard; DNA; 15 BP.
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- nsed

Norton JC;

Corey D,

Piatyszek MA,

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This is an oligonucleotide used in the measurement of Tm values and their
                                                                                                                                                                         New peptide nucleic acids hybridising to mammalian telomerase RNA - to inhibit telomerase, for treating tumours and other proliferative diseases, also for diagnosis.
                                                                                                                                                                                                                        Example 2; Page 49; 76pp; English.
                                                                       97WO-US005931.
                                                                                           96US-00630019
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (revised)
                                                                                                                                    Wright WE,
                                                                                                                                                         WPI; 1997-512647/47
                                                                                                               (GERO-) GERON CORP
                                                                                                                                                                                                                                                                                                                                                             Local Similarity
nes 15; Conserv
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modified_base
                             WO9738013-A1
                                                                       09-APR-1997;
                                                                                           09-APR-1996;
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12-MAY-1998
                                                  16-OCT-1997
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           Synthetic.
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                                                                                                                                     Shay JW,
                                                                                                                                                                                                                                                                                                                                                  Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                       This sequence is a novel peptide nucleic acid (PNA), which acts as an inhibitor of mammalian, preferably human, telomerase. The PNAs hybridise specifically to an RNA component of mammalian telomerase, and include the sequence GGG for appecific hybridisation to the template region of this component. PNAs can be used as probes to detect the RNA component of mammalian telomerage and as inhibitors of telomerase activity, especially
                                                                                                                                                                                                                                                                                                                                                            New peptide nucleic acids hybridising to mammalian telomerase RNA - used to inhibit telomerase, for treating tumours and other proliferative diseases, also for diagnosis.
                                                                                                                                                                             /note= "Sugar-phosphate backbone has been replaced by a peptide backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Revised record issued on 21-OCT-2004 : Correction to feature table key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Peptide nucleic acid; PNA; cancer; telomerase; probe; hybridisation; inhibitor; human telomerase RNA; hTR; PCR; oligonucleotide; ss.
                                                                                 Peptide nucleic acid; PNA; cancer; telomerase; probe; hybridisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA oligonucleotide 4, used in the measurement of Tm values.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 15; DB 1; Length 15;
Pred. No. 2.3e+02;
                                                            Peptide nucleic acid 5, targeted to mammalian telomerase.
                                                                                                                                                                                                                                                                                                                    Norton JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 15 BP; 5 A; 1 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                    Corey D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                    Piatyszek MA,
                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3.3%; Score
100.0%; Pre
0; h
                                                                                                                                                                    'mod base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                      Claim 9; Page 59; 76pp; English
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                                                                                                                                                                                                                                                      97WO-US005931
                                                                                                                                                                                                                                                                          96US-00630019
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         in the treatment of cancer
                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 100.
1es 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15 TTGTCTAACCCTAAC
                               (revised)
                                                                                                                                                                                                                                                                                                                    Wright WE,
                                                                                                                                                                                                                                                                                                                                         WPI; 1997-512647/47.
                                                                                                                                                                                                                                                                                                (GERO-) GERON CORP
                                                                                            inhibitor; ss.
                                                                                                                                     Key
modified_base
                                                                                                                                                                                                              WO9738013-A1.
                                                                                                                                                                                                                                                      09-APR-1997;
                                                                                                                                                                                                                                                                           09-APR-1996;
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                             21-OCT-2004
12-MAY-1998
                                                                                                                                                                                                                                  16-OCT-1997
                                                                                                               Synthetic
         AAT89229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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                                                                                                                                                                                                                                                                                                                    Shay JW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 297
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complimentary peptide nucleic acids (PNAs), (e.g. AAT89234). PNAs hybridise specifically to an RNA component of mammalian telomerase, and include the sequence GGG for specific hybridisation to the template region of this component. PNAs can be used as probes to detect the RNA component of mammalian telomerase and as inhibitors of telomerase activity, especially in the treatment of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /mod_base= OTHER
/note= "Sugar-phosphate backbone has been replaced by a
peptide backbone"
                                                                                                                                                                                                                                                                                                                 Gape
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Peptide nucleic acid; PNA; cancer; telomerase; probe; hybridisation; inhibitor; ss.
                                                                                                                                                                                                                                                                                                                 .;
0
                                                                                                                                                                                                                                                     3.3%; Score 15; DB 1; Length 15; 100.0%; Pred. No. 2.3e+02; ve 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Peptide nucleic acid 2, targeted to mammalian telomerase.
                                                                                                                                                                                                    Sequence 15 BP; 6 A; 4 C; 3 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
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                                                                                                                                                                                                                                                                                       100.08;
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                                                                                                                                                                                                                                                                                                                                                                                                  ACCCTAACTGAGAAG 15
                                                                                                                                                                                                                                                                                                                                                                     49 ACCCTAACTGAGAAG 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAT89226/c
ID AAT89226 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                 Conservative
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antisense oligonucleotides specifically hybridise to a nucleotide sequence within an accessible region of the hTR, but that does not sequence within an accessible region of the hTR, but that does not consider to a sequence within the template region of hTR. These oligonucleotides may specifically be used for detection of an RNA component of human telomerase in a sample. This is useful for diagnosing cancer (especially neuroblastoma, bladder, colon and prostate cancer), and providing prognosis for a cancer patient. The inhibitory oligonucleotides can inhibit the telomerase activity level in a cell by the transcription of the RNA component, decreasing the half-confidence of the component transcript, inhibiting assembly of the component into the telomerase holoenzyme, or inhibiting the polymerase activity of telomerase. These antisense oligonucleotides can be used for inhibiting telomerase activity in both cultured cells and in cells in vivo. They can be used in therapeutics for treating or preventing cancer, for contraception or sterilisation, for immunosuppression, and for selectively down-regulating specific branches of inflamma system, e.g. a specific subset of T-cells, in anti-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New peptide nucleic acid (PNA) compounds that inhibit telomerase activity in mammalian cells is useful as probes to detect the RNA component of a mammalian telomerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Peptide nucleic acid; PNA; telomerase; ribonucleoprotein enzyme; cancer; inhibitor; neoplasia; neurodegenerative disease; aging; hyperplasia; AIDS; HIV; fungal infection; forensic identification; detect; tumour; paternity testing; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "Peptide nucleic acid molecule, where N-(2-aminoethyl)glycine units are linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
oligonucleotides to the RNA component of human telomerase (hTR).
                                                                                                                                                                                                                                                                                                                                                                                                                                Score 15; DB 1; Length 15; Pred. No. 2.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PNA sequence #28 used to inhibit telomerase activity.
                                                                                                                                                                                                                                                                                                                                                                                            Sequence 15 BP; 5 A; 3 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Wright WE, Piatyszek MA, Shay JW, Norton JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 3.3%; Score 15; DB Best Local Similarity 100.0%; Pred. No. 2.3 Matches 15; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              152 CGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAA37570 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15 cerrcarreradade 1
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/*tag=
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                                                                                                                                                                                                                                                                                                                                                               parasites or fungi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-APR-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAA37570;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 300
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                임
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                         This sequence is a novel peptide nucleic acid (DNA), which acts as an inhibitor of mammalian, preferably human, telomerase. The PNAs hybridise specifically to an RNA component of mammalian telomerase, and include the sequence GGG for specific hybridisation to the template region of this component. PNAs can be used as probes to detect the RNA component of mammalian telomerase and as inhibitors of telomerase activity, especially
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         - used for detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           for treating cancers, contraception,
                                                                                                        New peptide nucleic acids hybridising to mammalian telomerase RNA - used to inhibit telomerase, for treating tumours and other proliferative diseases, also for diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RNA component; human telomerase; antisense oligonucleotide; infection; neuroblastoma; bladder cancer; colon cancer; prostate cancer; cancer; contraception; sterilisation; immunosuppression; therapeutic; hTR; immune system down-regulation; anti-inflammatory therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                   Revised record issued on 21-OCT-2004 : Correction to feature table key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RNA component of human telomerase (hTR) antisense oligo 21ab3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequences shown in AAV41169 to AAV41181 represent antisense
                                       Corey D, Norton JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New polynucleotide(s) anti:sense to human telomerase or inhibiting human telomerase, e.g. for treating car immuno-suppression or treating infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Weinrich SL;
                                                                                                                                                                                                                                                                                                                                                                                                                    Seguence 15 BP; 3 A; 2 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 15; DB 1; Le
Pred. No. 2.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pruzan R,
                                           Piatyszek MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 11; Page 65; 80pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                       7.34; Scc
Local Similarity 100.0%; Pr
nes 15; Conservative 0;
                                                                                                                                                                                            Claim 9; Page 59; 76pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 97WO-US023619
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96US-00770565
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     craacccraacreae 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAV41177 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                 the treatment of cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CTAACCCTAACTGAG
                                           Shay JW, Wright WE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1998-377670/32.
                                                                          WPI; 1997-512647/47.
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       (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAV41177;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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Corey DR;

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Gaps

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0; Indels

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The present sequence to the many component of the processe, and inhibits to telomerase activity. Telomerase is a ribonucleoprotein enzyme that synthesizes once strand of the telomeric DNA, using as a template an 11 nucleotide sequence contained within the RNA component of the enzyme. The nucleotide sequence contained within the RNA component of the enzyme. The invention relates to PNA molecules having a sequence of no more than 25 bases, which include the sequence GTTAGG. The uncharged nature of the PNA backbone increases the melting temperature of associating strands, increases the rate of association with targeted nucleic acids, and affords greater resistance of degradation by processes or nucleases. The therapeutic PNAs may be used for treating disease conditions such as cancers neoplasia, hyperplasia, neurodegenerative diseases, aging, human immunoseficiency virue (HIV) infection/AIDS (acquired immunodeficiency syndrome) and associated pathologies, fungal infections, and other cytooxic or diseases characterized by abnormal telomere metabolism or telomerase activity, in combination with antineoplastic and other cytooxic or cytostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization of individuals, e.g. paternity testing, based on hTR gene restriction of individuals, e.g. paternity testing, based on hTR gene restriction fragment length polymorphism (RPLP) pattern. PNAs are also useful as probes to detect the RNA component of amammalian telomerase and as allows cancerous conditions to be detected with increase and as allows cancerous conditions to be detected with increase and as and allows cancerous conditions to be detected with increase and as and allows cancerous conditions to be detected with increase and as and as a constitute.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             possibly at an earlier stage, before cells are detected as cancerous based on pathological characteristics. The diagnostic and prognostic methods of the present invention can be used to detect an immortal or neoplastic cell or tumour tissue or cancer of any oxigin, provided the cell expresses telomerase activity and its RNA component
                                                                    The present sequence represents a peptide nucleic acid molecule which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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/note= "Phosphorothioate internucleotide linkages"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Antisense sequence #45 used to inhibit telomerase activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3.3%; Score 15; DB 1; Length 15;
100.0%; Pred. No. 2.3e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 15 BP; 6 A; 4 C; 3 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                   Example 2; Col 33; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAA37587 standard; DNA; 15 BP
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/*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 301
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The present sequence represents an antisense oligonucleotide used as a control sequence alongside a peptide nucleic acid molecule which control sequence alongside a peptide nucleic acid molecule which control sequence alongside a peptide nucleic acid molecule which telomerase is a ribonucleoprotein enzyme that telomerase activity. Telomerase is a ribonucleoprotein enzyme that telomerase activity. Telomerase is a ribonucleoprotein enzyme that controlled the sequence contained within the RNA component of the enzyme. The invention relates to PNA molecules having a sequence of no more than 25 bases, which include the sequence GTTAGG. The uncharged nature of the PNA backbone increases the melting temperature of associating strands, increases the melting temperature of associating strands, increases the melting temperature of associating such as as concers, nooplasia, hyperplasia, neurodegenerative diseases or nucleases. The conclusion in an associated pathologies, fungal infections and other cycoxic or syndrome) and associated pathologies, fungal infections, and other conclusions, and other cycoxic or cytostatic agents, antifungal agents, and other nucleotise. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization probes to detect or quantitate polymucleotides having a human telomerase of individuals, e.g. paternity testing, based on hTR gene restriction of individuals, e.g. paternity testing, based on hTR gene restriction inhibitors of telomerase activity. The method of the present invention allows cancerous conditions to be detected with increased confidence and possibly at an earlier stage, before cells are detected as cancerous conditions to be detected with increased confidence and possibly at an earlier stage, before cells are detected as cancerous conditions to be detected and pathological characteristics. The diagnostic and propose to detect of control can be used as and sample and as a manally and the present invention allowers. The methods of the present invention are nearly an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Peptide nucleic acid; PNA; telomerase; ribonucleoprotein enzyme; cancer; inhibitor; neoplasia; neurodegenerative disease; aging; hyperplasia; AIDS; HIV; fungal infection; forensic identification; detect; tumour; paternity testing; ss.
                                                                                                                                                                                  New peptide nucleic acid (PNA) compounds that inhibit telomerase activilin mammalian cells is useful as probes to detect the RNA component of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                          Corey DR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3.3%; Score 15; DB 1; Length 15;
100.0%; Pred. No. 2.3e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cell expresses telomerase activity and its RNA component
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PNA sequence #2 used to inhibit telomerase activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 15 BP; 3 A; 2 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                          Norton JC,
                                                                                          Shay JW,
                                                                                                                                                                                                                                                                               Example 1; Col 27-28; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAA37545 standard; DNA; 15 BP
  96US-00630019.
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                                                                                          Piatyszek MA,
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Best Local Similarity 100.(
Matches 15; Conservative
                                            (TEXA ) UNIV TEXAS SYSTEM
                                                                                                                                                                                                                                      mammalian telomerase.
                                                                                                                                      WPI; 2000-292432/25
09-APR-1996;
                                                                                             Wright WE,
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AAA37545/c
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The present sequence to the many component of manmalian telomerase, and inhibits telomerase activity. Telomerase is a ribonucleoprotein enzyme that telomerase activity. Telomerase is a ribonucleoprotein enzyme that telomerase spatials as a template an 11 nucleotide sequence contained within the RNA component of the enzyme. The nucleotide sequence contained within the RNA component of the enzyme. The invention relates to PNA molecules having a sequence of no more than 25 bases, which include the sequence GTTAGG. The uncharged nature of the PNA backbone increases the melting temperature of associating strands, carfords greater resistance of degradation by proceases or nucleases. The therapeutic PNAs may be used for treating disease or nucleases. The therapeutic PNAs may be used for treating disease or nucleases. The cancers, neoplasia, hyperplasia, neurodegenerative diseases, aging, human confinction virus (HIV) infection/AIDS (acquired immunodeficiency syndrome) and associated pathologies, fungal infections, and other cytostatic agents, antifungal agents and other cytostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization cytostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization of individuals, e.g. paternity testing, based on hTR gene restriction fragment length polymorphism (RPLP) pattern. PNAs are also useful as probes to detect the RNA component of a mammalian telomerase and as inhibitors of telomerase activity. The method of the present invention continual conterns conditions to be detected with increase and as allows cancerous conditions to be detected with increase and as allows cancerous conditions to be detected with increase and as allows cancerous conditions to be detected with increase and as allows cancerous conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        wew peptide nucleic acid (PNA) compounds that inhibit telomerase activity in mammalian cells is useful as probes to detect the RNA component of a mammalian telomerase.
                                                                                             /note= "Peptide nucleic acid molecule, where N-(2-aminoethyl)glycine units are linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               possibly at an earlier stage, before cells are detected as cancerous based on pathological characteristics. The diagnostic and prognostic methods of the present invention can be used to detect an immortal or neoplastic cell or tumour tissue or cancer of any origin, provided the cell expresses telomerase activity and its RNA component
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    present sequence represents a peptide nucleic acid molecule which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Corey DR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Shay JW, Norton JC,
                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 6; Col 71; 45pp; English.
                                                                                                                                                                                                                                                                                                               97US-00838545.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Wright WE, Piatyszek MA,
                                                                                                                                                                                                                                                                                                                                                                                                                     (TEXA ) UNIV TEXAS SYSTEM
                                                                         *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2000-292432/25.
                           Key
misc feature
                                                                                                                                                                                                                                                                                                                  09-APR-1997;
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Sequence 15 BP; 3 A; 2 C; 5 G; 5 T; 0 U; 0 Other;

Query Match

Best Local Similarity 100.0%; Pred. No. 2.38+02;

Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps
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46 CTAACCCTAACTGAG 60 ||||||||||||||| 15 CTAACCCTAACTGAG 1

8 8

RESULT 303 AAA37548/c ID AAA37548 standard; DNA; 15

AAA37548;

15-AUG-2000 (first entry)

PNA sequence #5 used to inhibit telomerase activity.

Peptide nucleic acid; PNA; telomerase; ribonucleoprotein enzyme; cancer; inhibitor; neoplasia; neurodegenerative disease; aging; hyperplasia; AIDS; HIV; fungal infection; forensic identification; detect; tumour; paternity testing; 88.

Synthetic

Key Location/Qualifiers misc_feature 1..15

/*tag= a horizes - Peptide nucleic acid molecule, where N-(2-aninoethyl) glycine units are linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"

US6046307-A.

-APR-2000.

09-APR-1997; 97US-00838545.

09-APR-1996; 96US-00630019

(TEXA) UNIV TEXAS SYSTEM.

Wright WE, Piatyszek MA, Shay JW, Norton JC, Corey DR;

WPI; 2000-292432/25.

New peptide nucleic acid (PNA) compounds that inhibit telomerase activity in mammalian cells is useful as probes to detect the RNA component of a mammalian telomerase.

Claim 6; Col 71; 45pp; English.

The present sequence represents a peptide nucleic acid molecule which the present sequence represents a peptide nucleic acid molecule which component of mammalian telomerase, and inhibits telomerase activity. Telomerase is a ribonucleoprotein enzyme that stelomerase activity. Telomerase is a ribonucleoprotein enzyme that stelomerase activity. Telomerase is a ribonucleoprotein enzyme that component of the enzyme. The invention relates to PNA molecules having a sequence of no more than 25 hases, which include the sequence GTTAGG. The uncharged nature of the PNA backbon increases the melting temperature of associating strands, increases the rate of association with targeted nucleic acids, and affords greater resistance of degradation by proteases or nucleases. The therapeutic PNAS may be used for treating diseases or nucleases. The therapeutic PNAS may be used for treating diseases or nucleases. The therapeutic PNAS may be used for treating diseases or nucleases. The immunodeficiency virus (HIV) infection/AIDS (acquired diseases, adjng, human immunodeficiency virus (HIV) infection/AIDS (acquired immunodeficiency sydrome) and associated pathologies, fungal infections, and other cytotoxic or cytostatic agents, antitungal agents, and other nucleotides. PNAS may be used for molecular diagnostics, labelled PNAs are used as hybridization brobes to detect or quantitate polynucleotides having a human telomerase RNA (HTR) sequence. PNA probes are also used for forensic identification of individuals, e.g. paternity testing, based on hTR gene restriction fragment length polymorphism (RFLP) pattern. PNAs are also useful as chooses to detect the RNA component of a mammalian telomerase and as inhibitors of telomerase activity. The method of the present invention can be used to detected as cancerous based on pathological characteristics. The diagnostic and proposition cell expresses telomerase activity. The method of enceted and immortal or neoplastic cell or temorase doring the present invention cell expresses telomerase activi

Sequence 15 BP; 5 A; 1 C; 5 G; 4 T; 0 U; 0 Other;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to peptide nucleic acids (PNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in mammalian cells by hybridising to the RNA component of mammalian telomerase. The PNAs are useful as probes to detect the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polymucleotide having the human telomerase RNA component (hTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the hTR gene RFLP pattern. The PNA can be further used for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenetative diseases. The PNAs in combination with other pharmaceuticals (such as antineoplastic or cytostatic agents) can be used for treating neoplasia,
                                                                                                                                                                                                                                                                                                                                                                                  /*tag= a
/note= "This sequence is a peptide nucleic acid, i.e. it
contains a polyamide backbone instead of a deoxyribose
backbone"
                                  Gaps
                                                                                                                                                                                                                                   Mammalian; peptide nucleic acid; probe; forensic; paternity testing; human telomerase RNA component; hTR gene RFLP pattern; cancer; inflammation; lymphoproliferative disease; autoimmune disease; neurodegenerative disease; neoplasia; hyperplasia; HTV; AIDS; human immunodeficiency virus; acquired immunodeficiency syndrome; telomere metabolism; mutant; cytostatic; anti-inflammatory; immunosuppressive; polyamide backbone; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammetion, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases.
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                                                                                                                                                                                                               PNA XIII inhibiting human and mammalian telomerase activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Shay JW, Wright WE, Piatyszek MA, Corey DR, Norton JC;
            Length 15;
                                  0; Indels
           Score 15; DB 1; Le
Pred. No. 2.3e+02;
                                  Mismatches
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100.0%; Pred
0; M
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97US-00838545,
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AAS15427 standard; DNA; 15
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                                                                                                                                                                                       (first entry)
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                                  Conservative
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           Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                   modified base
                                                                                                                                                                                         14-FEB-2002
                                                                                                                                                                                                                                                                                                                                     sapiens
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09-APR-1997;
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                                                                                                                                                                                                                                                                                                                                                Synthetic.
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The present invention relates to peptide nucleic acids (PNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in mammalian cells by hybridising to the RNA component of mammalian cells by hybridising to the RNA component of mammalian cells by hybridising to the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polynucleotide having the human telomerase RNA component (hTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the hTR gene RFLP pattern. The PNA can be further used for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmaceuticals (such as antinooplastic or cytostatic agents) can be used for treating neoplasts, hyperplasia, human immunodeficiency virus (HIV) infections, acquired
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                                                                                                   activity. The present sequence represents one of the PNA sequences of the
hyperplasia, human immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomere metabolism or telomerase
                                                                                                                                                                                                                                                                                                                                  Gape
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2.3e+02;
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100.0%; Pred. No. 2.3
tive 0; Mismatches
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97US-00838545
                                                                                                                                                                                                      Sequence 15 BP; 5 A; 1 C; 5
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Best Local Similarity
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09-APR-1997;
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                                                                                                                                            invention
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/note= "This sequence is a peptide nucleic acid, i.e. it
contains a polyamide backbone instead of a deoxyribose
backbone"
          diseases characterised by abnormal telomere metabolism or telomerase activity. The present sequence representing a DNA oligonucleotide is complementary to some of the PNAs of the present invention, and is used
 immunodeficiency syndrome (AIDS) and associated pathologies, and other
                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                            Mammalian; peptide nucleic acid; probe; forensic; paternity testing; human telomerase RNA component; HTR gene RFLP pattern; cancer; inflammation; lymphoproliferative disease; autoimmune disease; neurodegenerative disease; neoplasia; hyperplasia; HIV; AIDS; human immunodeficiency virus; acquired immunodeficiency syndrome; telomere metabolism; mutant; cytostatic; anti-inflammatory; immunosuppressive; polyamide backbone; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, on neurodegenerative diseases.
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                                                                                           Length 15;
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                                                                   Sequence 15 BP; 6 A; 4 C; 3 G; 2 T; 0 U; 0 Other;
                                                                                          Score 15; DB 1; Le
Pred. No. 2.3e+02;
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                                                                                                      100.0%; Pred. No. 2.3
ive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
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                                              in melting temperature studies
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nes 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
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                                                                                            Query Match
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                                                                                                                                                                                                                                                                                                                       activity. The present sequence represents one of the PNA sequences of the
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component (hTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the hTR gene RPLP pattern. The PNA can be further used for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmaceuticals (such as antineoplastic or cytostatic agents) can be used for treating neoplasia, hyperplasia, human immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (ALIDS) and associated pathologies, and other diseases characterised by abnormal telemere metabolism or telemerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mammalian; forensic; paternity testing; human telomerase RNA component; hTR gene RFLP pattern; cancer; inflammation; lymphoproliferative diseas autoimmune disease; neurodegenerative disease; neoplasia; hyperplasia; HIV; AlDS; human immunodeficiency virus; telomere metabolism; mutant; acquired immunodeficiency syndrome; cytostatic; anti-inflammatory; immunosuppressive; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Phosphorothioate (PS) oligomer II used to inhibit telomerase activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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/label= OTHER
note= "Phosphorothioate internucleotide linkages"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 15; DB 1; Length 15;
Pred. No. 2.3e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 15 BP; 3 A; 2 C; 5 G; 5 T; 0 U; 0 Other;
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100.0%; Pred. No. ...
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Best Local Similarity 100.
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                     Invention
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a sequence of 6-25 nucleobases, that inhibit telomerase activity in mammalian cells by hybridising to the RNA component of mammalian telomerase. The PMAs are useful as probes to detect the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to
                                                                                 detect and/or quantitate polymicleotide having the human telomerase RNA component (HTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the hTR gene RFLP pattern. The PNA can be further used for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmaceuticals (such as antinoplastic or cytostatic agents) can be used for treating neoplasis hyperplasia, human immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomere metabolism or telomerase activity. The present sequence represents a phosphorothioate (PS) oligomer used to inhibit telomerase activity in the methods of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New polynucleotide useful for inhibiting telomerase activity in cells, of for treating telomerase-mediated condition or disease, such as cancers, tumors, Hodgkin's disease, or inflammatory conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma;
breast; prostate; colon; mixed cell leukaemia; Hodgkin's disease;
fertility; inflammatory condition; tumour; cancer; veterinary;
immunosuppression; telomerase inhibitor; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 15 BP; 3 A; 2 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human telomerase polynucleotide inhibitor #8.
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/*tag= a
/mod_base= OTHER
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ID AAS15927/C

XAC AAS15927 Standard; DNA; 15 BP.

AC AAS15927;

XX AAS15927;

XX T2-FEB-2002 (first entry)

XX Human; telomerase; hTR; cytostat

XM Human; telomerase; hTR; cytostat

XM Ertility; inflammatory condition

XM Munosuppression; telomerase in

XX Mosouphers.

XX Homo sapiens.

Synthetic.

XX Key I. 15

FT Modified_base I. 15

FT Modified_base I. 15

Yote= "N3'-P5'

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Best Local Similarity 100.
                                                                                                                                                                                                                                                                                                                                                                                    present invention
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The invention relates to polynucleotide inhibitors (I) and methods for inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity and proliferation of a telomerase posttive cell, and in manufacturing a medicament for inhibiting telomerase activity in a call manufacturing a medicament for inhibiting telomerase activity in a cell and in treating telomerase-mediated condition or disease, such as adenocarcinoma of breast, prostate or colon, mixed cell leukaemia, Hodgkin's disease, fertility and inflammatory conditions. (I) are also useful in treating a tumour or in manufacturing a medicament for the treatment of tumour. The polynucleotide inhibitors may also be used in diagnostic assays for detecting RNA or DNA. Inhibition of telomerase contivity in cells in vivo is useful in prophylactic and therapeutic methods of treating cancer and other disorders involving inappropriate expression of telomerase, and in treating veterinary proliferative continuation of telomerase in haematopoietic stem cells is useful for immunosuppression and for selectively down-regulating specific continuation system. The present sequence represents human telomerase polynucleotide inhibitor #8, as described in the method of the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma;
breast; prostate; colon; mixed cell leukaemia; Hodgkin's disease;
fertility; inflammatory condition; tumour; cancer; veterinary;
immunosuppression; telomerase inhibitor; ss.
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100.0%; Pred. No. 2.3e+02;
/ative 0; Mismatches 0; Indels
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/note= "N3'-P5' phosphoramidate linkages"
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AAS15931 standard; DNA; 15
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Best Local Similarity
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The invention relates to polymucleotide inhibitors (I) and methods for inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity and proliferation of a telomerase positive cell, and in manufacturing a medicament for inhibiting telomerase activity in a cell can in treating telomerase-mediated condition or disease, such as adenocarcinoma of breast, prostate or colon, mixed cell leukaemia, CK Gadenocarcinoma of breast, prostate or colon, mixed cell leukaemia, CK Hodgkin's disease, fertility and inflammatory conditions. (I) are also useful in treating a tumour or in manufacturing a medicament for the ctreatment of tumour. The polymucleotide inhibitors may also be used in catement of tumour or in useful in prophylactic and therapeutic activity in cells in vivo is useful in prophylactic and therapeutic methods of treating cancer and other disorders involving inappropriate expression of telomerase, and in treating veterinary proliferative configences in thibition of telomerase in haematopoletic stem cells is useful for immunosuppression and for selectively down-regulating specific branches of the immune system. The present sequence represents human the immense plynucleotide inhibitor #12, as described in the method of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma; breast; prostate; colon; mixed cell leukaemia; Hodgkin's disease; fertility, inflammatory condition; tumour; cancer; veterinary; immunosuppression; telomerase inhibitor; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 3.3%; Score 15; DB 1; Length 15; Best Local Similarity 100.0%; Pred. No. 2.3e+02; Matches 15; Conservative 0; Mismatches 0; Indels
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Example 3; Page 32; 48pp; English.
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/*tag= a
/mod_base= OTHER
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AAS15932 standard; DNA; 15 BP.
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                                                                                                                                                                                                                                                                                                                                                                         the invention
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modified_base
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The invention relates to polymucleotide inhibitors (I) and methods for inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity and proliferation of a telomerase positive cell, and in manufacturing a medicament for inhibiting telomerase activity in a cell and in treating telomerase-mediated condition or disease, such as adenocarcinoma of breast, prostate or colon, mixed cell leukaemia. (Consection of disease, fertility and inflammatory conditions. (I) are also treatment of tumourr or in manufacturing a medicament for the treatment of tumourr or in manufacturing a medicament for the creatment of tumourr. The polymucleotide inhibitions and therapeutic diagnostic assays for detecting RNA or DNA. Inhibition of telomerase cation disorders involving inappropriate expression of telomerase, and in treating veterinary proliferative capression of telomerase, and in treating veterinary proliferative for immunosuppression and for selectively down-regulating specific branches of the immune system. The present sequence represents human the inversion that in the present sequence represents human the inversion.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                             3.3%; Score 15; DB 1; Length 15;
100.0%; Pred. No. 2.3e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                         Sequence 15 BP; 5 A; 1 C; 7 G; 2 T; 0 U; 0 Other;
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/mod_base= OTHER
/note= "A-hydroxyethyl phosphate"
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              Example 3; Page 32; 48pp; English.
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/mod_base= OTHER
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Best Local Similarity 100.
Matches 15, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Koizumi M, Morita K;
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modified_base
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12-OCT-1998
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                                                                                                                                                                        The present invention relates to novel 2',5'-oligoadenylic acid analogs and their pharmacologically- acceptable salts. The analogs are stable with superior antitumour and antiviral activity and so are useful in cancer or antiviral therapy e.g. as antisense drugs. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          note = "2',4'-oxyethylene linkage in the sugar residues"
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nucleic acid units as well as unusual phosphate groups with excellent activity particularly antitumor, applicable in cancer or antiviral
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/note= "A-hydroxyethyl phosphate"
                                                                                                                  Disclosure; Page 100; 220pp; Japanese
                                                                                                                                                                                                                                                                                       was used to illustrate the invention.
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Best Local Similarity 100.
Matches 15; Conservative
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                                                            therapy
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                                                                 The present invention relates to novel 2',5'-oligoadenylic acid analogs and their pharmacologically- acceptable salts. The analogs are stable with superior antitumour and antiviral activity and so are useful in cancer or antiviral therapy eg. as antisense drugs. The present sequence was used to illustrate the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                3.3%; Score 15; DB 1; Length 15; 100.0%; Pred. No. 2.3e+02; Ve 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                           Sequence 15 BP; 5 A; 3 C; 7 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human telomerase antisense RNA primer.
Disclosure; Page 100; 220pp; Japanese.
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97US-00873039.
97US-00951733.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        80 TITIGCICCCGCGC 94
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(first entry)
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CANADA INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense compounds targeted to nucleic acid molecules encoding acyl-
coenzyme A synthetase 1 (ACSI), useful for treating diseases or
conditions associated with aberrant expression of ACSI, e.g. diabetes,
                                                                            Gaps
                                                                                                                                                                                                                                                          Acyl-coenzyme A synthetase 1, ACS1, antisense oligonucleotide #205.
                                                                                                                                                                                                                                                                                 acyl-coenzyme A synthetase 1; ACS1; diabetes; obesity;
metabolic syndrome X; cardiovascular disorder; cancer; infection;
inflammation; tumour; antisense; ss.
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                                             Ouery Match 3.3%; Score 15; DB 1; Length 18; Best Local Similarity 100.0%; Pred. No. 2.9e+02; Matches 15; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3.3%; Score 15; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02; ive 0; Mismatches 0; Indels
                      Sequence 18 BP; 1 A; 5 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 3 A; 12 C; 2 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 3; SEQ ID NO 205; 940pp; English.

    ACS1, antisense oligonucleotide.

                                                                                                                                                                                  ADK20128 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                          14-AUG-2003; 2003WO-US025389.
                                                                                                                                                                                                                                                                                                                                                                                                                                    14-AUG-2002; 2002US-0403591P.
                                                                                                              GGCTGGCCTGGGAG 18
                                                                                               11 GGGTGGGCCTGGGAG 25
                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                        (PHAA ) PHARMACIA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2004-203782/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       obesity or cancer.
                                                                                                                                                                                                                                                                                                                                                          WO2004016749-A2
                                                                                                                                                                                                                                  18-NOV-2004
                                                                                                                                                                                                                                                                                                                                                                                  26-FEB-2004
                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                           ADK20128;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ROSS SA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
field.)
                                                                                                                                                            RESULT 314
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The invention relates to an antisense compound targeted to a nucleic acid molecule encoding acyl-coenzyme A synthetase 1 (ACS1). The antisense compound specifically hybridises with and inhibits the expression of ACS1. The antisense oligonucleotides or compounds are useful for inhibiting the expression of acyl-coenzyme A synthetase 1 (ACS1), and for treating diseases or conditions associated with aberrant expression of ACS1, e.g. diabetes, obesity, metabolic syndrome X, cardiovascular disorder or cancer. The antisense compounds are also useful as research reagents and kits, or in diagnostic, therapeutic and prophylactic applications, e.g. to prevent or delay infection, inflammation or tumour formation. The present sequence represents an acyl-coenzyme A synthetase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New antisense compounds targeted to nucleic acid molecules encoding acyl-coenzyme A synthetase 1 (ACS1), useful for treating diseases or conditions associated with aberrant expression of ACS1, e.g. diabetes,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cold-adapted influenza virus; passage culture; PB2 protein; PB1 protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                             Acyl-coenzyme A synthetase 1, ACS1, antisense oligonucleotide #221.
                                                                                                                                                                                                                    acyl-coenzyme A synthetase 1; AC31; diabetes; obesity;
metabolic syndrome X; cardiovascular disorder; cancer; infection;
inflammation; tumour; antisense; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 3 A; 13 C; 2 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Influenza virus PA gene specific primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 3; SEQ ID NO 221; 940pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1, ACS1, antisense oligonucleotide
ADK20144 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14-AUG-2003; 2003WO-US025389.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14-AUG-2002; 2002US-0403591P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           335 GGGCGAGGCGAGGT 349
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match

Query Match

Best Local Similarity 100.00

"Thes 15; Conservative
                                                                                                        18-NOV-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAX82232 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20 GGGCGAGGCCGAGGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-203782/19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 obesity or cancer.
                                                                                                                                                                                                                                                                                                                                                                                        WO2004016749-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                               26-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18-AUG-1999
                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                    ADK20144;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ross SA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAX82232;
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AAX82232/c
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Gaps

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GGGCGAGGCGAGGT 349

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Local Similarity 100. 1es 15; Conservative

Best Loca Matches

19 GGGCGAGGCGAGGT 5

99WO-US008731 98US-0082477P

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A novel rapid and sensitive method using PCR for detecting Histoplasma
                                                                                            (USSH ) US DEPT HEALTH & HUMAN SERVICES.
                                                                                                                                                                                       capsulatum in samples.
                                                                                                                     Schafer MP, Reid TM;
                                                                                                                                              WPI; 2000-052703/04.
                                       20-APR-1999;
                                                                 21-APR-1998;
              28-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAQ23265;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     셤
                                                                                                                                                                                                                                                                                                                                   The invention relates to cold-adapted influenza viruses prepared by passage culture of A/X-31, B/Yamagata/16/88 or B/Leo/40 viruses at low temperatures. A cDNA gene of cold-adapted influenza virus HTCA-A101 can be selected from a group consisting of PB2 protein gene, PB1 protein gene, RAPR2122-X82197). The method is useful for the production of coldadapted influenza virus that exhibit temperature sensitivity and can be actively grown in fertilized eggs. The virus is useful for vaccines for protection against fil. Live vaccines containing cold-adapted viruses have several advantages over killed vaccines. It can prevent reduction of immunogenicity, which may occur in the killed vaccine where antigenic protecins will be denatured at its inactivation. It can also avoid typersensitivity due to the prolonged administration of heterologous proteins. It promotes the immunity by inducing IgA and it can be administrated into a spray formulation via nasal cavity and thus its administration is convenient for children. It able to inhibit the growth of the wild-type virus and thus its therapeutic effect can be expected. Sequences AXX8222-X82257 represent PCR primers specific for the various
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
PA protein, NP protein, M protein, NS protein, temperature sensitivity, vaccine, flu, influenza, PCR primer, ss.
                                                                                                                                                                                                                                                                    Cold-adapted influenza viruses useful for the production of protective
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                  Kim HG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5.8S rRNA, detection, environmental sample, public health, soil, vegetation; dust, faeces, histoplasmosis, PCR primer; ss.
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 14.8; DB 1; Length 18;
Pred. No. 3e+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                  Kim J,
                                                                                                                                                                                                                  Cheoun KH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 18 BP; 8 A; 4 C; 5 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      H. capsulatum 5.8S rRNA gene PCR primer HC2.
                                                                                                                                                                                                                  Kim SJ,
                                                                                                                                                                                                                                                                                                              Example 4; Page 15; 62pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29 TGGTGGCCATTTTTGTC 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18 recreecearricare 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAZ24498 standard; DNA; 18 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3.3%;
                                                                                                                                   98WO-KR000384.
                                                                                                                                                                                                                  Youn JW,
                                                                                                                                                              97KR-00064854
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(first entry)
                                                                                                                                                                                        (CHEI-) CHEIL JEDANG CORP.
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Best Local Similarity 88.9,
Best Local Similarity
The Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    genes of influenza virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ajellomyces capsulatus
                                                                                                                                                                                                                                                                                     vaccines against flu.
                                                                                                                                                                                                                  Lee KH,
                                                                                                                                                                                                                                           WPI; 1999-385377/32.
                                                     Influenza virus.
                                                                                                                                   30-NOV-1998;
                                                                                                                                                              29-NOV-1997;
                                                                              WO9928445-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9954508-A1
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18-FEB-2000
                                                                                                         10-JUN-1999
                                        Synthetic.
                                                                                                                                                                                                                   Seong BL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAZ24498;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 317
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This invention describes a novel rapid and sensitive method for detecting Histoplasma capsulatum. The method uses oligonuclectide primers which amplify a segment of DNA specific to the H. capsulatum 5.85 rRNA gene. The method of the invention can be used to detect H. capsulatum in a wide variety of samples. For example, for public health protection convironmental samples containing soil, vegetation, dust, decaying faeces from birds and bats, etc. can be assayed for the presence of the canalysed e.g. those who are suspected of having contracted fresh histoplasmosis as a result of symptomology, and immune compromised individuals who may be at risk of contracting ofhonic pulmonary or ocular chaptasmosis. The prior arisk of contracting ofhonic pulmonary or ocular chaptasmosis. The prior arisk of contracting ofhonic pulmonary or capsulatum is expensive and requires several weeks to complete. The capsulatum is expensive and requires several weeks to complete. The capsulatum is expensive and requires several weeks complete. The capsulatum. The present invention provides this needs as it is rapid, sensitive and inexpensive. This sequence represents PCR primer HCI which is used in the method of detection. (Updated on 15-SEP-2003 to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3.3%; Score 14.8; DB 1; Length 18; 88.9%; Pred. No. 3e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 18 BP; 3 A; 8 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Japanese type C hepatitis virus partial gene.
Claim 2; Page 30; 33pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 410 CTGAGCTGTGGGACGTGC 427
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAQ23265 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HCV; probe; diagnostic; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (TOKU ) TOKUYAMA SODA KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13-JUN-1990;
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                                                                                                   The polynucleotide contains the partial base sequence of the Japanese type C hepatitis virus (HCV) gene. The DNA was obtd. from the serum of a non A non B hepatitis patient. The serum was ultracentrifuged and the ppte. dissolved in a mixture of GITC, Na citrate, sarcosine and mercaptoethanol. Phenol chloroform was added and the mixture centrifuged. Isopropanol was added to the aq. layer which was left to stand at -20 deg C for 3 hrs, then centrifuged. The ppte. was treated by GITC and phenolchloroform and pptd. By ethanol and dissolved in dR2O to give an aq. DNA soln. The gene was sequence A genomic version of the sequence was given (AAQ23036). In the version shown there is a Y at base 189. The DNA in the specification has a guanine at this position. The DNA may be used as a probe for detection of Japanese type C hepatitis. See also AAQ230264
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention describes a method for inducing apoptosis in a cell or animal comprising administering to a cell or animal a Jagged 2
                                Poly:nucleotide for hepatitis virus gene detection - using poly:nucleotide deoxyribonucleic acid probe, hybridising with hepatitis virus gene and detecting hybrid for diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       apoptosis; Jagged 2 inhibitor; cytostatic; hyperproliferative disorder; human; ss; PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Inducing apoptosis in a cell or animal for treating a subject having a condition associated with insufficient apoptosis by administering to a cell or animal a Jagged 2 inhibitor to reduce Jagged 2 levels or
                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                 3.2%; Score 14.6; DB 1; Length 18; 82.4%; Pred. No. 3.18+02; tive 2; Mismatches 1; Indels
                                                                                                                                                                                                                                                                          Sequence 18 BP; 1 A; 10 C; 5 G; 0 T; 0 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human Jagged 2 forward PCR primer SEQ ID NO:4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 13; SEQ ID NO 4; 148pp; English
                                                                                  Disclosure, Page 2; 14pp, Japanese.
                                                                                                                                                                                                                                                                                                                                                 196 CGCCCTCCCGGGGACC 212
                                                                                                                                                                                                                                                                                                                                                              *GCCCCCCYGGGGACC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                  BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12-MAR-2002; 2002US-00096399
                                                                                                                                                                                                                                                                                                                                                                                                                                  ADD00949 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                          14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Shapard PJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-756943/71.
           WPI; 1992-101941/13.
                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO2003077848-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Koller E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADD00949;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          activity.
                                                                                                                                                                                                                                                                                                  Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antisense, Jagged 2; hyperproliferative disorder; cancer; developmental disorder; apoptosis; prophylaxis; antisense therapy; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense oligonucleotides for modulating Jagged 2 expression, useful for diagnosing, preventing or treating diseases or conditions associated
                    treating a subject having a disease or condition associated with insufficient apoptosis by administration of a Jagged 2 inhibitor; (2) a pharmaceutical composition comprising a Jagged 2 inhibitor and another active ingredient for inducing apoptosis; and (3) a kit comprising a Jagged 2 inhibitor and instructions for using the Jagged 2 inhibitor in the induction of apoptosis. The Jagged 2 inhibitor has cytostatic activity. The method can be used for inducing apoptosis in a call or animal for treating a subject having a disease or condition associated with insufficient apoptosis, e.g., hyperproliferative disorder. The present sequence represents a PCR primer for human Jagged 2, which is used in an example from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
inhibitor to reduce Jagged 2 levels or activity. Also described: (1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          for diagnosing, preventing or treating diseases or con with Jagged 2, e.g. cancer or developmental disorders
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 16 BP; 1 A; 7 C; 5 G; 3 T; 0 U; 0 Other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3.2%; Score 14.4; DB 1;
Local Similarity 93.8%; Pred. No. 2.8e+02;
les 15; Conservative 0; Mismatches 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  265 CCCGGGGCTTCTCCGG 280
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-898250/82.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 primer; PCR; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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Matches
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G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; oesophageal cancer; scancer; black cancer; band cancer; band and neck cancer; ovarian cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxel; docetaxel; cisplatin; methotrexxate; gemcitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatoid arthritis; restenosis; Crohm's disease; obesity; sechsemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsie; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammatory bowel disease; infection; se.

Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme;

NFKB sub-unit modulating inozyme substrate #146.

(first entry)

03-JUN-2003

ACA06327;

BP.

ACA06327 standard; RNA; 17

ACA06327 ID ACA(

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This invention relates to novel antisense compounds that can be used to modulate the expression of Jagged 2. Specifically, it refers to compositions useful for inhibiting the expression of Jagged 2, a human homologue of the Drosophila Scrate gene, which is involved in differentiation and cell fate, as well as positive feedback control over signalling genes such as Notch 1, Notch 3 and Jagged 1. The Jagged 2 gene is located on chromosome 14932, a region that has been implicated in genetic diseases including Usher syndrome type la that is associated with cetinitis pigmentosa. The present invention describes antisense oligonucleotides that comprise at least one modified sugar moiety, a 2'-O methoxyethyl (2' MOE) and at least one modified nucleobase, a 5-C methoxyethyl (2' MOE) and at least one modified nucleobase, a 5-C methylcytosine. These compounds are useful for modulating the mucleic acid molecules encoding Jagged 2, ultimately modulating the amount of Jagged 2 produced, which in turn is useful for research
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New antisense oligonucleotides of 8-40 nucleobases, useful for modulating the function of nucleic acid molecules encoding Jagged 2, ultimately modulating the amount of Jagged 2 produced.
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                                                                                                                                                                                                                                                                                                                                                                  human, Jagged 2; differentiation; cell fate; signalling;
Usher syndrome type la; retinitis pigmentosa; PCR; ss; primer.
3.2%; Score 14.4; DB 1; Length 16; 93.8%; Pred. No. 2.8e+02; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   primer used to amplify human Jagged 2 DNA of the invention
                                                                                                                                                                                                                                                                                                                              PCR primer used to amplify human Jagged 2 DNA SeqID 4.
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                                                                              280
                                                                                                                                                                                                          ADH57064 standard; DNA; 16 BP
                                                                                                  1 CCCAGGGCTTCTCCGG 16
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                                                                              265 CCCGGGGCTTCTCCGG
                                                                                                                                                                                                                                                                                        (first entry)
Query Match 3.2
Best Local Similarity 93.8
Matches 15; Conservative
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                    Best Loca
Matches
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92US-00987132. 94US-00245466. 94US-00291932. 96US-00777916.

STINCHCOMB D T. MCSWIGGEN J.

(STIN/)

(DRAP/) DRAPER K G.

23-MAY-2001; 2001US-00864785

07-DEC-1992; 18-MAY-1994; 15-AUG-1994; 23-DEC-1996;

Homo sapiens. JS2002177568-A1

28-NOV-2002

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Stinchcomb DT, Mcswiggen J, Draper KG;

WPI; 2003-340953/32.

Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.

Claim 3; Page 29; 72pp; English.

The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NRKB), where I) is an inozyme, zinzyme, G-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulatings REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A.

(I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, espocially Mg 2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer. malanoma, lymphoma, glioma or multidurg resistant cancer antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, edatrexate, genetitable or radiation therapy. The enzymatic and antisense nucleic acid molecules are also useful for treating inflammatory disease such as calle useful for treating inflammatory disease such as calle useful for treating inflammatory disease, diabetes, obesity, autoimmune disease, lupus, multiple solerosis, transplant/graft rejection, gene therapy applications, ischemialreperfusion injury (central nervous system (CNS) and molecules of a novel enzymatic en ensymatic central nervous system (GNS) and molecules of a novel enzymatic.

infection. This sequence represents the substrate of a novel enzymatic
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Gaps

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3.2%; Score 14.4; DB 1; Length 16; 93.8%; Pred. No. 2.8e+02; ve 0; Mismatches 1; Indels

llarity 93.8%; Conservative 0

Best Local Similarity Matches 15; Conserv

Query Match

265 CCCGGGGCTTCTCCGG 280

16

CCCAGGCTTCTCCGG

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RESULT 322

SXS

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BLATT L.
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(MACE/)
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(I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially Mg<sup>2</sup>2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                              Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; dung cancer; brain cancer; brain cancer; prostate cancer; colorectal cancer; prain cancer; loosesophageal cancer; stomach cancer; bladder cancer; pancreatic cancer; cervical cancer; had and neck cancer; ovarian cancer; melanoma; lymphoma; gloma; multidrug resistant cancer; REL-A-specific inhibitor; commontherapy; paclitaxal; docetaxel; cisplatin; methotrexate; commonterapy; paclitaxal; fluorouracil carboplatin; edatrexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatorid arthritis; restencés; Crohn's disease; obesity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammation; inflammatory bowel disease; infection; ss.
                                                                                                              Gaps
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                                                                       3.2%; Score 14.4; DB 1; Length 17; 75.0%; Pred. No. 3e+02; tive 3; Mismatches 1; Indels
                                     Sequence 17 BP; 0 A; 10 C; 4 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                     NFKB sub-unit modulating inozyme substrate #145.
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94US-00291932.
96US-00777916.
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                                                                                            Best_Local Similarity 75.0
Matches 12; Conservative
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MCSWIGGEN J.
DRAPER K G.
   nucleic acid molecule
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23-DEC-1996;
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                                                                           Query Match
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prostate, colorectal, brain, oesophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A-specific inhibitors or chemocherapy including paclitaxel, docetaxel, cisplatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, gencitabine or radiation therapy. The enzymatic and antisense nucleic acid molecules are also useful for treating inflammatory disease such as theumatoid arthritis, restenosis, asthma, Crohn's disease, diabetes, obesity, autoimmune disease, lughs, multiple sclarosis, transplant/graft rejection, gene therapy applications, ischaemia/reperfusion injury (central nervous system (CNS) and myocardial), glomerulonephritis, espsis, allergic alrway inflammation, inflammatory bowel disease or infection. This sequence represents the substrate of a novel enzymatic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to an enzymatic nucleic acid molecule which specifically cleaves RNA derived from hepatitis C virus (HCV), in which the binding arms of the enzymatic nucleic acid molecule comprises sequences complementary to any of the defined substrate sequences given in the specification. The nucleic acid molecule may be administered for the treatment of HCV infections, especially in combination with type I interferons. The present sequence represents a HCV DNAzyme substrate
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HCV infection; type I interferon; DNAzyme.
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Best Local Similarity 75.0
Matches 12, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nucleic acid molecule
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MACEJACK D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hepatitis C virus.
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Unidentified,
                              24-FEB-2003
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                                                                                                                                                   19-SEP-2002
                                                                                                                                                                                                                                     Gillies SD,
                                                                                                                                                                                                                                                                                                               Ig moiety
          AAD47637;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAF57369;
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence is an antisense oligonucleotide which was used in an RNase mapping experiment. This enables the identification of sites within the Ii RNA strand which hybridise to antisense DNA. These sites can then be used as targets for antisense strands which may, using gene therapy, be used as tumour cell vaccines (for example to treat carcinomas, paranoman, laukaemia, lymphomas, stomach, breast, colon or rectum, lung, prostate, bladder, pancreas, brain and ovarian cancers), or they can be used to treat autoimmune diseases including rheumatoid arthritis,
                                                                                                                                                                                                                                                Immunoregulator; antisense oligonucleotide; cancer; tumour cell vaccine; rheumatoid arthritis; autoimmune disease; diabetes mellitus; thyroiditis;
                                                                                                                                                                                                                            Antisense oligonucleotide for use in RNase H mapping assay SEQ ID NO: 24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cancer cell vaccine for treating malignancies, autoimmune disorders and isolating autodeterminant peptides comprises a regulator of invariant chain protein expression or immunoregulatory function.
                                                             Gaps
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                                        Query Match 3.2%; Score 14.4; DB 1; Length 17; Best Local Similarity 82.4%; Pred. No. 3e+02; Matches 14; Conservative 1; Mismatches 2; Indels
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                    Sequence 17 BP; 5 A; 6 C; 4 G; 0 T; 1 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 18 BP; 3 A; 7 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 46; 94pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       diabetes mellitus and thyroiditis
                                                                                   432 AGGACTCGGCTCACACA 448
                                                                                                       1 AGGACUNGGCCCACACA 17
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                                                                                                                                                                                                                                                                                                                                                                                                                         Xu M, Qiu G, Humphreys R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Local 15; Conservative
These 15; Conservative
                                                                                                                                                             AAA63120 standard; DNA; 18
                                                                                                                                                                                                      (first entry)
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                                                                                                                                                                                  AAA63120;
  Bequence
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AAD47637/c
ID AAD476
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                                                                                                                                                           Immunoglobulin; Ig; antibody; immunocytokine; immunofusion; immunoligand;
protease resistance; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             we fusion protein, useful for producing antibodies, immunocytokines or tusion proteins that enhance the expression or protesse resistance of fusion protein, comprises an immunoglobulin (Ig) moiety fused to a non-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cdc25; Cdc25 phosphatase; transcription; modulator; murine; CDc25A; exon; intron; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           present invention relates to methods and compositions for efficiently
                                                                           Forward primer, to construct human KSgamma2h antibody expression vector.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Murine Cdc25A intron 7/exon 8 splice junction sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 5; Page 63; 71pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           300 GAAGAGTTGGGCTCTG 315
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(first entry)
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nes 15; Conserv
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The present invention relates to detecting and identifying fungal pathogenic species in a sample. The method involves hybridizing a nucleic acid of a fungal pathogen possibly present in the sample with at least one oligonucleotide probe, from an Internal Transcribed Spacer (ITS) region. The method is useful for simultaneous detection and differentiation of clinically important fungi in a single assay, Cartuelly indicatent fungi in a single assay. Cartuell, C. glabrata, C. parapsilosis, C. tropicalis, C. ketyr, versicole, A. nidulans, A. funigatus, C. neoformans and pneumocystis carinii. The method is especially useful in the detection of opportunistic infections in patients with impaired immunity systems, such as organ transplants patients, patients receiving intensive anticancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The patent discloses new macromolecules of formula PNA-DNA-PNA, in which DNA comprises at least one 2'-deoxynucleotide and each PNA comprises at least one peptide nucleic acid subunit. These compounds have increased resistance to nuclease and increased specific binding affinity, and they can activate RNaseH for target strand cleavage. They can hybridise specifically to a nucleic acid strand (especially RNA) and are useful (1) for treating diseases associated with undesirable production of protein,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New chimeric macromolecules contg. DNA and peptide nucleic acid segments - with good nuclease stability and binding affinity, also activating RNaseH, useful for treating disease, in diagnosis and for identifying
Cryptococcus and Aspergillus, comprises hybridizing the amplified nucleacid of the fungal pathogen with a probe from the internal transcribed spacer region of a DNA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oligonucleotide used in PNA-DNA-PNA chimeric macromolecule.
                                                                                                                                                                                                                                                                                                                                                                                                          DB 1; Length 17; 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                                                                                                                                                                       Sequence 17 BP; 1 A; 7 C; 7 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                          3.1%; Score 14; DB 100.0%; Pred. No. 3.3 tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                       treatments, diabetics or AIDS patients
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                                                                               Claim 1; Page 46; 59pp; English.
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Best Local Similarity 100.
Matches 14; Conservative
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           셤
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                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a method of identifying a modulator of Cdc25 activity that comprises contacting a test cell having a recombinant Cdc25 phosphatase gene whose expression alters transcription of a selected gene, with a compound under conditions where recombinant Cdc25 phosphatase gene is expressed and alters the transcription of a selected gene as an indication of the compound being a modulator of Cdc25-mediated transcription. The method is useful for identifying modulators of Cdc25
                                                                                                                                                                                                                                                             Assay for screening modulators of Cdc25 activity by using a cell having recombinant Cdc25 phosphatase gene whose expression alters the transcription of a selected gene in the presence of a modulator.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    activity. Sequences AAF57363-376 represent intron/exon splice junction sequences of the murine Cdc25A gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Detecting and identifying fungal pathogens, especially Candida
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14; DB 1; Length 17;
Pred. No. 3.3e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fungal pathogenic species identification probe #21.
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(IRBI-) ENTERPRISE IRELAND T/A BIORESEARCH IRELA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Jannes G,
                                                                                                                                                                                                                                                                                                                                         Example 1; Page 15; 55pp; English
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                                                                                                                         99US-0153639P
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                                                                                     11-SEP-2000; 2000WO-US024838
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Best Local Similarity
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                                                                                                                                                           (BADI ) BASF AG
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               WO200120034-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unidentified
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11-JUN-1999;
                                                                                                                         13-SEP-1999;
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                                                   22-MAR-2001
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                                                                                                                                                                                               Voss J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Fungal
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes an enzymatic nucleic acid molecule (I) with RNA cleaving activity, which modulates the expression of a plant gene. Also described is a gene comprising a cDN sequence encoding maize Delta-9 desaturase. (I) can be used to modulate expression of a gene, preferably Delta-9 desaturase or a granule bound starch synthase (GBSS) gene, in a plant (preferably a maize or canola plant). (I) can be used to modulate caffeine synthesis in a coffee plant, nicotine production in a tobacco plant, fruit ripening processes in an apple, tomato, pear, plum or peach plant, flower pigmentation in a rose, petunia, chrysanthemum or marigold plant or lignin production in a tobacco, aspen, poplar or pine
            for
          contacting a test solution with the macromolecule and RNaseH), or (3) fo in-vivo enhancement of polynucleotide hybridisation and RNase activity. They can also be used diagnostically and for screening chemotherapeutic
                                                                                                                                                                                                                                                                                                                                              Maize; corn; Zea mays; delta-9 desaturase; GBSS; target; substrate; granule bound starch synthase; hammerhead ribozyme; hairpin ribozyme; modulation; gene expression; transgenic plant; cleavage; canola plant; caffeine synthesis; coffee plant; nicotine production; tobacco; fruit ripening; flower pigmentation; lignin production; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Skokut TA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ribozyme which modulates plant gene expression - preferably modulates expression of DELTA-9 desaturase or granule bound starch synthase in maize or canola.
                                                                                                                            Gaps
in-vitro modification of sequence-specific nucleic acid (by
                                                                                                                            ö
                                                                                                 3.1%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 3.4e+02; ive 0; Mismatches 2; Indel8
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                                                                                                                                                                                                                                                                                                                       Delta-9 desaturase hamerhead ribozyme target SEQ ID NO:826.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Guo L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Merlo PAO,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17 BP; 1 A; 6 C; 5 G; 0 T; 5 U; 0 Other;
                                                                         Sequence 17 BP; 2 A; 5 C; 3 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mcswiggen JA,
Merlo DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 38; Page 86; 155pp; English.
                                                                                                                                                     102 TICTCGCTGACTITCAG 118
                                                                                                                                                                            1 rrcrcccrccarrrcag 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                            AAX62951 standard; RNA; 17
                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                            15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Edington BE,
Folkerts O,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (DOWC ) DOWELANCO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-JUL-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Zwick MG,
                                                                                                                                                                                                                                                                                                                                                                                                                              Зеа шаув.
                                                                                                                                                                                                                                                                     AAX62951;
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                                                                                                                          Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New nucleic acid(s) for regulating the Grb2-related with Insert Domain (GRID) gene comprises using antisense and enzymatic nucleic acid
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                                                                                                                                                                                                                                                                                                                                                           Human, Grb2-related with Insert Domain, GRID, T-cell;
co-stimulatory adaptor protein; tissue rejection, graft rejection;
leukaemia; cytostatic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ellis JH;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 13.8; DB 1; Length 17;
Pred. No. 3.4e+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hamblin PA,
                                                                                                                                                                                                                                                                                                                Human GRID NCH ribozyme substrate oligonucleotide #151
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mcswiggen JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    molecules such as hammerhead ribozymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 4; Page 65; 108pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         used to illustrate the invention
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106 CGCTGACTTTCAGCGGG 122
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                                             cecueccurucaecues
                                                                                                                                                                  ABL46697 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (RIBO-) RIBOZYME PHARM INC. (GLAX ) GLAXO GROUP LTD.
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                                                                                                                                                                                                                                                                                                                                                                                                                   leukaemia; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-550088/61.
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Best Local Similarity
Matches 15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
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                                                                                                                                                                                                                     ABL46697;
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ABL46496/C
ID ABL4649
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AC ABL464
XX
DT 27-JUN
XX
XX
DE Human
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Randi AM;

Von Carlowitz I, Mcswiggen JA, Mclaughlin F,

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Novel polynucleotide which down regulates expression of Ets-related gen useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome.
                                                                                                                                                                                                                         Claim 4; Page 125; 149pp; English.
                                       16-MAY-2000; 2000US-00572021.
             16-MAY-2001; 2001WO-US015866.
                                                                  (RIBO-) RIBOZYME PHARM (GLAX ) GLAXO GROUP LTD
                                                                                                                                     WPI; 2002-082995/11.
                                                                                                            Jarvis T,
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                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to oligonucleotides that downregulate the expression of human Grb2-related with Insert Domain (GRID) gene. GRID is a T-cell co-stimulatory adaptor protein. The oligonucleotides are useful for modulating the expression of GRID, to treat conditions such as tissue/graft rejection and leukaemia. The oligonucleotides can also be administered in conjunction with other therapies such as radiation, chemotherapy and cyclosporin treatment. The present oligonucleotide was used to illustrate the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; hammerhead ribozyme; cytostatic; antitumour; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; virucide; osteopathic; vulnerary; cancer; lymphoma; Ewing's sarcoma; melanoma; psoriasis; tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgaris; angiofibroma of tuberous solerosis; port-wine stain; wound healing; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leukaemia; ss; Osler-Weber-rendu syndrome, leukaemia; osteoporosis; DNAzyme; inozyme;
                                                                                                                                                                                                                                                                                                           New nucleic acid(s) for regulating the Grb2-related with Insert Domain (GRID) gene comprises using antisense and enzymatic nucleic acid molecules such as hammerhead ribozymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
        Human, Grb2-related with Insert Domain, GRID; T-cell; co-stimulatory adaptor protein; tissue rejection; graft rejection; leukaemia; cytostatic; ss.
                                                                                                                                                                                                                                                      Ellis JH;
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                                                                                                                                                                                                                                                      Mcswiggen JA, Hamblin PA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17 BP; 3 A; 5 C; 8 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                    Claim 4; Page 61; 108pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            201 CTCCCGGGGACCTGCGG 217
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                                                                                                                                                     23-FEB-2001; 2001WO-US005957.
                                                                                                                                                                                 24-FEB-2000; 2000US-0184594P.
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                                                                                                                                                                                                          (RIBO-) RIBOZYME PHARM INC.
(GLAX ) GLAXO GROUP LTD.
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nes 15; Conservative
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                                                                                               WO200162911-A2.
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                                                                    Homo sapiens
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                                                                                                                         30-AUG-2001.
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ABK19302/c
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The invention relates to a nucleic acid molecule (I) which down regulates expression of an Ets-related gene (ERG). (I) is useful for treating conditions selected from cancer, lymphome, Ewing's sarcoma, metanoma, conditions selected from cancer, lymphome, Ewing's sarcoma, metanoma, tumour angiogenesis, diabetic retinopathy, macular degeneration, tumour angiogenesis, diabetic retinopathy, macular degeneration, verruca vulgaris, angiofibroma of tuberous sclerosis, port wine stains, Sturge Neber syndrome, Rippel Trenaunay-Meber syndrome, Oaler-Weber-rendu Syndrome, leukaemia, osteoporosis and wound healing. (I) is useful for treating a patient having a condition associated with the level of ERG, by contacting cells of the patient with (I) under conditions suitable for the treatment. Leukaemia or tumour angiogenesis is treated by administering (I) to the patient in conjunction with one or more of other therapies such as radiation or conjunction with one or more of other therapies such as radiation or chemotherapy treatment. (I) is useful for reducing ERG activity in a cell, by contacting the cell with (I). (I) is useful for cleaving RNA of ERG gene, by contacting (I) with RNA, in the presence of a divalent cellseases related to the expression of ERG, and as diagnostic tool to examine genetic drift and mutations within diseased cells or to detect the presence of ERG RNA in a cell. (I) is useful for specifically transcring molecules which regulate expression of ERG fusion genes. ABX 1354-ABX2719 represent nucleic acids, including antisense and enzymetic nucleic acids molecules which regulate expression of ERG, and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 13.8; DB 1;
Pred. No. 3.4e+02;
0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   related PCR primers of the invention
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Best Local Similarity 88.2
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                     proteins and their coding sequences: MD23, MD24, MD27, MD212. MD23 is proteins and their coding sequences: MD24, MD24, MD27, MD212. MD23. MD212. MD21 is encoded at chromosome fg21.1, MD24 is encoded at chromosome fg21.3-22.2, MD24. MD23, MD24, MD27, and MD212 sequences are useful in therapy. Or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MD23, MD24, MD27, or MD212, e.g. cancer or developmental disorders. The nucleic caused by altered expression of MD23, MD24, MD27, or MD212. The nucleic acids can also be used as probes to detect and characterize gross alterations in MD23, MD27, or MD212. The nucleic acids can also be used as probes to detect and characterize gross alterations in MD23, MD27, or MD212 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymetic nucleic acid; hammer/head ribozyme; DNAzyme; inozyme; inozyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative; disease state, HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
                                                                                                                                                                                        New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                           The present invention relates to novel human zinc finger-containing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.1%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 3.48+02; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17 BP; 2 A; 2 C; 8 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                               Example 8; SEQ ID NO 814; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HCV DNAzyme substrate sequence #800.
                                                                                                                                                                                                                                  MDZ4, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24 AGGGGTGGTGGCCATTT 40
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                                         30-JUL-2002; 2002EP-00016874
                                                                      02-AUG-2001; 2001US-00922181
                                                                                                                              Gu Y, Nguyen C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15; Conservative
                                                                                                                                                            WPI; 2003-423107/40.
                                                                                                  (AEOM-) AEOMICA INC
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Best Local Similarity
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             05-FEB-2003
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                                                                                                                               Shannon M,
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the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, Indxymes, indxymes, and G-claever ribozymes, DNAzymes, or nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptase and/or HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV DNA. The nucleic acids may be used to modulate the expression of HBV genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and complete the expression and/or replication of HCV. The compounds and disease states related to HBV and HCV infection, replication and gene cates related to HBV and HCV infection, replication and gene active acids and complete the expression such as cirrhosis, liver failure, and hepatocellular carcinoma. The present sequence represente a substrate for one of the HCV DNAzyme or minus strand DNAzyme sequences disclosed in the present
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to nucleic acid molecules which modulate
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                                                                                                                                                                                                                                                                                                                                                                                                                                Lee
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                                                                                                                                                                                                                                                                                                                                                                                                                                Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HCV minus strand DNAzyme substrate sequence #853.
                                                                                                                                                                                                                                                                                                                                                                                                                           Mcswiggen J, Morrissey D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 17 BP; 5 A; 6 C; 5 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 248; 387pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    432 AGGACTCGGCTCACACA 448
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08-JUN-2001; 2001US-00877478.
08-JUN-2001; 2001US-0296876P.
24-OCT-2001; 2001US-0335659P.
05-DEC-2001; 2001US-0337055P.
                                                                                                                                           RIBOZYME PHARM INC.
BLATT L.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Roberts E;
                                                                                                                                                                                             MACEJAK D.
MCSWIGGEN J.
MORRISSEY D.
PAVCO P.
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                                                                                                                                                                                                                                                                                                                                      DRAPER K.
ROBERTS E.
                                                                                                                                                                                                                                                                                                                  LEE P.
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(DRAP/)
(ROBE/)
                                                                                                                                           (RIBO-)
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Matches
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Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; murine; tumour suppression; tumour reversion; apoptosis; virus resistance; viral disease; tumour; cell degeneration; cancer; Alzheimer's disease;

schizophrenia; ss.

#02003025176-A2. Mus musculus,

27-MAR-2003

Murine oligonucleotide associated with tumour supression, SEQ ID 5862.

01-JUL-2003 (first entry)

ACC68615;

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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, inozymes, zinzymes, amberzymes, anderzymes, and have viral replication, Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and disease states related to HBV and HCV infection, replication and gene expression such as cirrhosis, liver failure, and hepatocallular expression such as cirrhosis, liver failure, and hepatocallular expression and NNAzyme sequences disclosed in the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
 degenerative; disease state; HBV infection; HCV infection; cirrhosis;
liver failure; hepatocellular carcinoma; hepatotropic; cytostatic;
virucide; antiinflammatory; substrate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pavco P, Lee P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mcswiggen J, Morrissey D,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 290; 387pp; English.
                                                                                                                                                                                                        26-MAR-2001; 2001US-00817879.
08-UIN-2001; 2001US-00877478.
08-UIN-2001; 2001US-0296876P.
24-CCT-2001; 2001US-0335659P.
05-DEC-2001; 2001US-0337055P.
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                                                                                                                                                                                                                                                                                                            RIBOZYME PHARM INC.
BLATT L.
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Roberts E;
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                                                                                                                                                                                                                                                                                                                                              MACEJAK D.
                                                                     Hepatitis C virus.
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LEE P.
DRAPER K.
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                                                                                                       WO200281494-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    invention
                                                                                                                                                                                                                                                                                                                         (BLAT/)
(MACE/)
(MCSW/)
(MORR/)
(PAVC/)
(LEEP/)
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                                                                                                                                                                                                                                                                                                            (RIBO-)
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New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies

Tuijnder

Amson R,

Telerman A,

WPI; 2003-333167/31.

(MOLE-) MOLECULAR ENGINES LAB.

17-SEP-2002; 2002WO-IB004210. 17-SEP-2001; 2001FR-00011979. Disclosure; Page 716; 738pp; French.

and transfected cells.

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The present invention relates to murine oligonucleotides (ACC62754-ACC68006), which are associated with tumour suppression, tumour reversion, apoptosis and virus resistance. The oligonucleotides are useful as (1) as probes and primers for detecting, identifying, quantifying and/or amplifying uncleic acid, e.g. as one component of a gene chip; in vitro as (anti) eense reagents; and (2) for production of recombinant polypeptides. The oligonucleotides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restenosis; asthma; Crohn's disease; diabbtes; obesity; autoimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human NOGO receptor zinzyme substrate sequence #84.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 1 A; 6 C; 2 G; 8 T; 0 U; 0 Other;
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tes 15; Conservative
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Gaps

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3.1%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 3.4e+02; 1ve 0; Mismatches 2; Indels

88.2%;

Conservative

Local Similarity les 15; Conserv

Matches

Query Match

TGGGAGGGGTGGCC 36

20 17

recenecarectic 1

BP.

ACC68615 standard; DNA; 17

RESULT 337 ACC68615/c ID ACC686 XX

asthma; allergic rhinitis; atopic dermatitis; human PTGDR;

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allergy; as
substrate;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention comprises nucleic acids (e.g. antisense oligonucleotides)
that down regulate the expression or inhibit the function of a receptor
for a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (PTGDR),
IKappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the
invention are useful for treating: cerebrovascular accident, central
nervous system (CNS) injury, spinal cord injury, cancer (e.g. melanoma,
lymphoma or glioma), inflammatory disease (e.g. rheumatoid archritis,
restenosis or asthma), inflammatory disease (e.g. rheumatoid archritis,
cestenosis or asthma), crohn's disease, diabetes, obesity, autoimmune
ischaemia/reperfusion injury, glomerulonephritis, sepsis, and allergic
conditions (e.g. asthma, allergic rhinitis or atopic dermatitis). The
nucleic acids of the invention are also useful for down-regulating the
cexpression of a target gene and as a diagnostic tool to examine genetic
drifts and mutations within diseased cells or to detect the presence of a
target RNA in a cell. The present RNA sequence represents a human NOGO
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                                                                                                                                                                                                                                                                                                                                                                                               Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restenosis; asthma; Crohn's disease; diabetes; obesity; autoimmune disease; lupus; multiple sclerosis; transplant rejection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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Pred. No. 3.4e+02;
2; Mismatches 2; Indels
allergy, asthma; allergic rhinitis; atopic dermatitis; NOGO receptor zinzyme; substrate; ds.
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11 Similarity 76.5%;
13; Conservative 2
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28-AUG-2001; 2001US-0315315P.
                                                                                                                                                                        03-APR-2002; 2002WO-US010512
                                                                                                                                                                                                           05-APR-2001; 2001US-00827395
                                                                                                                                                                                                                                                                                     (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                                                                                                                                          Blatt L, Chowrira B,
                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-058513/05.
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Best Local Similarity
                                                                                               WO200281628-A2
                                                          Unidentified
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ID ADL51
XX
AC ADL51
XX
XX
DT 20-M
XX
XX
KW anti
KW pros
KW pros
KW pros
KW mela
KW mela
KW rest
KW auto
KW graf
KW auto
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The invention comprises nucleic acids (e.g. antisense oligonucleotides)
that down regulate the expression or inhibit the function of a receptor
for a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (FTGDR),
IKappa Kinase (IKK), or protein Kinase PKR. The nucleic acids of the
invention are useful for treating: cerebrovascular accident, central
nervous system (CNS) injury, spinal cord injury, cancer (e.g. melanoma,
nervous system (CNS) injury, spinal cord injury, cancer (e.g. melanoma,
restenosis or asthma), Crohn's disease, diabetes, obesity, autoimmune
disease, lugus, multiple sclerosis, transplant/graft rejection,
cischaemia/reperfusion injury, glomerulonephritis, sepsis, and allergic
conditions (e.g. asthma, allergic rhinitis or atopic dermatitis). The
nucleic acids of the invention are also useful for down-regulating the
cxpression of a target gene and as a diagnostic tool to examine genetic
drifts and mutations within diseased cells or to detect the presence of
target RNA in a cell. The present RNA sequence represents a human PKR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; prostaglandin D2 recebrovascular accident; central kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restenoisis; asthma; Crohn's disease; diabbtes; obesity; autoimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Fosnaugh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17 BP; 9 A; 2 C; 5 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Haeberli P, Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 161; SEQ ID NO 5454; 317pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human IKK-gamma substrate sequence #814.
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                                                                                                                                                                                                                                                                                             05-APR-2001; 2001US-00827395.
29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
                                                                                                                                                                                                                              03-APR-2002; 2002WO-US010512
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Best Local Similarity 88.20,
                                                                                                                                                                                                                                                                                                                                                                                                                                                        (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Blatt L, Chowrira B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-058513/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        substrate sequence.
                                                                           WO200281628-A2
Unidentified
                                                                                                                                                  17-0CT-2002
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The invention comprises nucleic acids (e.g. antisense oligonucleotides)
that down regulate the expression or inhibit the function of a receptor
for a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (PTGDR),
IRappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the
invention are useful for treating: creebrovascular accident, central
nervous system (CKS) injury, spinal cord injury, cancer (e.g. melanoma,
Impubna or glioma), inflammatory disease (e.g. rheumatoid arthritis,
restenosis or asthma), Crohn's disease, disease, obesity, autoimmune
disease, lupus, multiple sclerosis, transplant/graft rejection,
conditions (e.g. asthma, allergic rhinitis or atopic dermatitis). The
nucleic acids of the invention are also useful for down-regulating the
conditions (e.g. asthma, allergic rhinitis or atopic dermatitis). The
nucleic acids of the invention are also useful for down-regulating the
carget session of a target gene and as a diagnostic tool to examine genetic
drifts and mutations within diseased cells or to detect the presence of a
target RNA in a cell. The present RNA sequence represents a human IKK-
gamma substrate sequence.
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  allergy; asthma; allergic rhinitis; atopic dermatitis; Human IKK-gamma;
                                                                                                                                                                                                                                                                                                                                                                                                                     Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antisense oligonucleotide, neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; ecrebrovascular accident; protein kinase PKR; ecrebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restenosis; asthma; Crohn's disease; diabetes; obesity; autorimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17 BP; 1 A; 7 C; 7 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 59; SEQ ID NO 1837; 317pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human IKK-gamma substrate sequence #813.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            249 TGGAGGCCGCGGTCGGC 265
                                                                                                                                                                                                                      05-APR-2001; 2001US-00827395.
29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                  WO200281628-A2
                         substrate; ds
                                                              Unidentified.
                                                                                                                                          17-0CT-2002
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The invention comprises nucleic acids (e.g. antisense oligonucleotides)
that down regulate the expression or inhibit the function of a receptor
for a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (PrGDR),
KappaB kinase (IKK), or protein kinase PrR. The nucleic acids of the
invention are useful for treating: cerebrovascular accident, central
nervous system (CNS) injury, spinal cord injury, cancer (e.g. melanoma,
restenosis or asthma), inflammatory disease, diabetes, obesity, autoimmune
classes, lupus, multiple sclerosis, transplant/graft rejection,
ischaemia/reperfusion injury, glomerulonephritis, sepsis, and allergic
conditions (e.g. asthma, allergic rhinitis or atopic dermattis). The
nucleic acids of the invention are also useful for down-regulating the
expression of a target gene and as a diagnostic tool to examine genetic
drifts and mutations within diseased cells or to detect the presence of
carget evaluations within diseased cells or to detect the presence of
carget evaluations within diseased cells or to detect the presence of
carget evaluations within diseased cells or to detect the presence of
carget evaluations within diseased cells or to detect the presence of
carget evaluations.
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allergy; asthma; allergic rhinitis; atopic dermatitis; Human IKK-gamma;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; ss; GRID; Grb2-related with insert domain; hammerhead ribozyme;
NCH ribozyme; G-cleaver ribozyme; Zinzyme; DNAzyme; amberzyme; Inozyme;
halrpin ribozyme; tissue rejection; graft rejection; leukaemia.
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                                                                                                                                                                                                                                                                                                                                  Haeberli P, Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 17 BP; 1 A; 6 C; 8 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 59; SEQ ID NO 1836; 317pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human GRID mRNA substrate sequence #129.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              262
                                                                                                                                                                                                             05-APR-2001; 2001US-00827395.
29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADM53854 standard; mRNA; 17 BP
                                                                                                                                                                           03-APR-2002; 2002WO-US010512,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 GCGUGGAGGCCGCCCC 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   76.5%;
                                                                                                                                                                                                                                                                                         (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gamma substrate sequence.
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                                                                                                                                                                                                                                                                                                                                Chowrira B,
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                                                                                               WO200281628-A2.
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                    substrate; ds.
                                                         Unidentified.
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                                                                                                                                    17-0CT-2002
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Matches
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leukemia
                                    (JARV/)
                                                                                   ELLI/)
                                                 (CARL/)
(MCSW/)
                                                                         HAMB,
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                                                                                                                                                                                                                                                                                The invention relates to a nucleic acid molecule that down-regulates expression of Grb2-related with insert domain (GRID) gene, e.g. a hammerhead ribozyme, NCH ribozyme, G-cleaver ribozyme, Zinzyme, DNAzyme, amberzyme, Inozyme or hairpin ribozyme. Also include are a mammalian cell including the novel nucleic acid molecule, reducing GRID activity in a cell by contacting the cell with the novel nucleic acid molecule, treating a patient having a condition associated with the level of GRID creating a patient having a condition associated with the level of GRID cell or treating a patient having a condition associated with the level of GRID contacting the cell with the novel nucleic acid molecule, cleaving RNA of a GRID gene by contacting the cell with the novel nucleic acid molecule in ammaner that allows its expression vector comprising a nucleic acid sequences (encoding at least the novel nucleic acid molecule in ammaner that allows its expression), a mammalian cell including the expression vector and an enzymatic nucleic acid molecule is useful for treating a condition associated with the level of GRID, e.g. tissue/graft rejection and leukaemia. The present sequence is a target region for the enzymatic nucleic acide of the invention.
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                                                                                                                                                                                              New nucleic acid molecule that down-regulates expression of Grb2-related with insert domain (GRID) gene, useful for treating a condition associated with the level of GRID, e.g. tissue/graft rejection and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; 88; GRID; Grb2-related with insert domain; hammerhead ribozyme;
NCH ribozyme; G-cleaver ribozyme; Zinzyme; DNAzyme; amberzyme; Inozyme;
hairpin ribozyme; tissue rejection; graft rejection; leukaemia.
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                                                                                                                                                Ellis JH
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                                                                                                                                                Mcswiggen J, Hamblin PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17 BP; 3 A; 5 C; 8 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human GRID mRNA substrate sequence #330.
                                                                                                                                                                                                                                                            Claim 4; SEQ ID NO 129; 74pp; English.
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                       23-FEB-2001; 2001US-00792818
                                               10-FEB-2000; 2000US-0181594P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       201 CTCCCGGGGACCTGCGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                Jarvis T, Carlowitz IV,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity 88.2
les 15; Conservative
                                                                      ) JARVIS T.
CARLOWITZ I V.
MCSWIGGEN J.
HAMBLIN P A.
                                                                                 (CARL/) CARLOWITZ I (MCSW/) MCSWIGGEN J (HAMB/) HAMBLIN P A (ELLI/) ELLIS J H.
                                                                                                                                                                        WPI; 2003-829646/77
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17-JUL-2003
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                                                                                                                                                                                                                                     leukemia.
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                                                                      (JARV/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 343
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The invention relates to a nucleic acid molecule that down-regulates expression of Grb2-related with insert domain (GRID) gene, e.g. a hammerhead ribozyme, NCH ribozyme, G-cleaver ribozyme, Zinzywe, DNAzyme, C amberzyme, Inozyme or hairpin ribozyme. Also include are a mammalian cell molecule including the novel nucleic acid molecule, reducing GRID activity in a cell by contacting the cell with the novel nucleic acid molecule, treating a patient having a condition associated with the level of GRID c treating a patient having a condition associated with the level of GRID c contacting the cell with the novel nucleic acid molecule, cleaving RNA of a GRID gene by contacting the cell with the novel nucleic acid molecule, an expression vector comprising a nucleic acid sequences (encoding at less the novel nucleic acid molecule in a manner that allows its expression), a mammalian cell including the expression vector and an enzymatic nucleic coid molecule that cleaves RNA derived from a GRID gene. The nucleic acid molecule that cleaves RNA derived from a GRID gene. The nucleic acid molecule that cleaves RNA derived from a GRID gene. The nucleic acid molecule that cleaves RNA derived from a GRID gene. The nucleic acid molecule that cleaves RNA derived from a GRID gene. The nucleic acid molecule is useful for treating a condition associated with the level of GRID, e.g. tissue/graft rejection and leukaemia. The present sequence is a target region for the enzymatic nucleic acide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New nucleic acid molecule that down-regulates expression of Grb2-related with insert domain (GRID) gene, useful for treating a condition associated with the level of GRID, e.g. tissue/graft rejection and
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                                                                                                                                                                                                                                                                                                                                             Ellis JH;
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                                                                                                                                                                                                                                                                                                                                             Hamblin PA,
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                                                                                                                                                                                                                                                                                                                                             Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 4; SEQ ID NO 330; 74pp; English.
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10-FEB-2000; 2000US-0181594P.
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ID ADI85884 standard; RNA; 17
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                                                                                                                                                                                                                                                                                                                                             Carlowitz IV,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                     CARLOWITZ I V. MCSWIGGEN J. HAMBLIN P A. ELLIS J H.
                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-829646/77.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US2003125270-A1
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                                                                                                                                                                                                                                                                                                                                             Jarvis T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AD185884;
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Claim 42; Paqe 83; 155pp; English.
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                                                                                                                   Enzymatic nucleic acid molecules which specifically cleave RNA derived from hepatitis C virus (HCV), useful for the treatment of HCV infections, especially in combination with type I interferon therapy.
                                                                                                                                                                                                The invention relates to an enzymatic nucleic acid molecule which specifically cleaves RNA derived from hepatitis C virus (HCV), in which the binding arms of the enzymatic nucleic acid molecule comprises sequences complementary to any of the defined substrate sequences given in the specification. The nucleic acid molecule may be administered for the treatment of HCV infections, especially in combination with type I interferons. The present sequence represents a HCV DNAzyme substrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Maize, corn, Zea mays, delta-9 desaturase, GBSS, target, substrate, granule bound starch synthase; hammerhead ribozyme, hairpin ribozyme, modulation; gene expression, transgenic plant; cleavage; canola plant; caffeine synthesis; coffee plant; nicotine production; tobacco; fruit ripening; flower pigmentation; lignin production; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Skokut TA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ribozyme which modulates plant gene expression - preferably modulates expression of DELTA-9 desaturase or granule bound starch synthase in
                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Granule bound starch synthase hairpin substrate SEQ ID NO:571.
                                                                                                                                                                                                                                                                                                                                                3.1%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 3.48+02; ive 0; Mismatches 2; Indels
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                                                                 Pavco PA, Macejack
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Merlo DJ;
                                                                                                                                                                                                                                                                                                                       Sequence 17 BP; 3 A; 11 C; 2 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                          Claim 1; SEQ ID NO 3130; 198pp; English
                                                                   Roberts E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP
                                                                                                                                                                                                                                                                                                                                                                                                       TGGGAGGGGTGGTGGCC 36
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15; Conservative
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Folkerts O,
                                                                   Blatt L, Mcswiggen J,
 MCSWIGGEN J.
             ROBERTS E.
PAVCO P A.
MACEJACK D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1997-202224/18.
                                                                                            WPI; 2004-031273/03
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Young SA,
                                                                                                                                                                                                                                                                                                 sequence.
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                          (PAVC/)
(MACE/)
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 (MCSM/)
               ROBE/)
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maize or canola.

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The present invention describes an enzymatic nucleic acid molecule (I) with RNA cleaving activity, which modulates the expression of a plant gene. Also described is a gene comprising a cDNA sequence encoding maize Delta-9 desaturase. (I) can be used to modulate expression of a gene, preferably Delta-9 desaturase or a granule bound starch synthase (GBSS) gene, in a plant (preferably a maize or canola plant). (I) can be used to modulate caffeine synthesis in a coffee plant, nicotine production in a tobacco plant, fruit ripening processes in an apple, tomato, pear, plum or peach plant, flower pigmentation in a rose, petunia, chrysanthemum or marigold plant or lignin production in a tobacco, aspen, poplar or pine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This exon specific primer used in the PCR amplification of a short stature associated sequence. The gene region corresponding to short stature has been identified as a region of approximately 500 kb in the PARI region of the X and Y chromosomes. There genes in this region have been identified as candidates for the short stature gene. These genes were designated SHOX (also referred to as SHOX93 or HOX93), pET92 and SHOX (slox-like homeobox gene on chromosome three). The SHOX gene has two separate splicing sites resulting in two variations SHOXa and SHOXb. The specification provides sequences of SHOX (short stature homeobox-containing) genes SHOX ET92, SHOXb, SHOT and exons of the SHOX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homeobox domain; human growth gene; growth regulation; growth defect; turner's syndrome; short stature homeobox containing gene; short stature; SHOX; bone disease; osteoporosis; calcium regulation; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New human growth genes - used to develop products for the diagnosis and treatment of human growth defects such as short stature, e.g. Turner's
                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                               Sequence 18 BP; 1 A; 9 C; 7 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                    1; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                              134 CGGCCTGCCGCCTTCCA 150
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97EP-00100583.
                                                                                                                                                                                                                                                                                                           3.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ä
                                                                                                                                                                                                                                                                                                                                                                                                                                         CGGCCUGCCGCCGGCCA
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Best Local Similarity 82.4
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1998-271719/24
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AAV35627/c
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genes as shown in AAV35610 to AAV35621 and protein sequences of the human growth protein transcription factor SHOXa, SHOXb and SHOT as shown AAW60573. The novel genes are responsible for human growth. Defects in the genes acan cause short stature, e.g. Turner's syndrome. The products can be used to develop agents for the treatment of short stature providing a mitogenic effect on cells, e.g. for the treatment of bone diseases such as osteoporosis and diseases involved with disturbance in the bone calcium regulation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Probe 52 used to detect polymorphism in human HLA-DRB1 exon 2 DNA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   ch 3.1%; Score 13.8; DB 1; Length 18; 1 Similarity 88.2%; Pred. No. 3.6e+02; 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                  Sequence 18 BP; 0 A; 14 C; 4 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            332 CGGGGGCGAGGCGAGG 348
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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Best Local Similarity
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Synthetic.
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This invention relates to a novel chemical sensor system method. Specifically, it refers to an isolated nucleic acid molecule that encodes a receptor protein, which binds to chemicals that can stimulate the sense of taste or smell for example. The present invention describes the manufacture of a chip that acts as support to immobilise transfected calls expressing the receptor gene, such that this chip can be employed as a component of the chemical sensor model. Furthermore, this chip is useable as an artificial sensory organ where the chemical receptor occurains an olfactory receptor the sensor can react to olfactory stimulation. Accordingly, these sensors are useful in the food industry for analysing freshness of meat, fruit and vegetables, hygiene simple city manufaction, environmental examination and disease diagnosis. Purthermore, such systems are automatable for high throughput applications under various conditions, even for differentiating optical isomers of R(:)-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chemical sensor systems based on chemical receptors introduced into cells for immobilization onto support to form chip as component of sensor, useful in detecting stimuli e.g. taste and smell applicable in food
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 carvone from S(+)-carvone easily. This oligonucleotide sequence is used to construct an odourant receptor DNA expression vector of the invention
                                                                                                                                                                                                                                             DNA oligo to construct an odourant receptor expression vector SegID102.
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                                                                                                                                                                                                                                                                              ss; chemical sensor system; taste; smell; artificial sensory organ; olfactory stimulation; food industry; hygiene inspection; environmental examination; disease diagnosis; carvone.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Yoshikawa T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Seguence 18 BP; 5 A; 8 C; 4 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (NAAD-) NAT INST ADVANCED IND SCI & TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 9; SEQ ID NO 102; 521pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sato T, Hirono J, Hamana H, Miyake M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         237 CCGAACCCCGCCTGGAG 253
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-MAY-2002; 2002JP-00154239.
13-JUN-2002; 2002JP-00172412.
14-JAN-2003; 2003JP-00005175.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-MAY-2003; 2003WO-JP006719
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 88.2%;
262 CGGCCGGGGCTTCTCC
                           CGGCCCGGGCCTGCTCC
                                                                                                                              ADM42858 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ccgaacaccccrccac
                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-023356/02.
                                                                                                                                                                                                                                                                                                                                                                                                   WO2003100057-A1
                                                                                                                                                                                                         03-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                        04-DEC-2003
                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                     18
                                                                                                                                                                     ADM42858;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     industry.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                            RESULT 348
                                                                                                                ADM42858
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RESULT 349

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Gaps

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3.1%; Score 13.8; DB 1; Length 18; 88.2%; Pred. No. 3.6e+02; ive 0; Mismatches 2; Indels

Conservative

Query Match Best Local Similarity Matches 15; Conserva

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27-JUN-2003 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The INVELLOR interacts to promoter regions from micros and indicated to a cancer, especially a gene encoding a cytotoxin, for therapy of the terologous gene, especially a gene encoding a cytotoxin, for therapy of cancer, especially neoplasias. The telomerase is necessary for the unrestricted proliferative capacity of many human cancers. Mutation or dysregulation of telomerase repression pathway may cause reactivation or upregulation of telomerase expression in cancer. Substances, identified in the methods, can be used to block transcription from the TR gene promoter through interaction of the 5' regulatory sequences. These substances, e.g. antisense oligonucleotides, transcription factors, peptide nucleic acids and factors that disrupt signal transduction, are useful for cancer therapy. In particular, gene therapy vectors (especially pGT62-codAupp) comprising the promoter and a viral thymidine kinase gene can be used to convert a prodrug, e.g. gancyclovir, so that neoplasia can be controlled or treated. Direct down-regulation of telomerase RNA gene through manipulation of transcription factors may be effective anticancer therapy and the cloning of the HTR gene promoter allows the analysis of therapeutic molecules which modulate hTR promoter activity. Sequences AAZ07696-3121 represent PCR primers used in cloning and mutagenists of human TR gene (HTR) promoter region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ·.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to promoter regions from mouse and human telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                 Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; gene therapy; thymidine kinase gene; anticancer therapy; human; mutagenesis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ;
0
                                                                                                                                                                                                 Human telomerase RNA gene (hTR) promoter specific primer hll.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 13.8; DB 1; Length 38; Pred. No. 4.8e+02; 0; Mismatches 7; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 38 BP; 2 A; 7 C; 23 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CANC-) CANCER RES CAMPAIGN TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    222 TCGCCTGCCCAGCCCCCGAACCCCG 246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    rcccaegcccaccrccccacaccc
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                                  BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99WO-GB000308
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Best Local Similarity 72.0%;
                               AAZ07296 standard; DNA; 38
                                                                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                           22-OCT-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gene therapy
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                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Keith WN;
                                                                                     AAZ07296;
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AAZ07296/c
ID AAZ072
XX
AC AAZ072
XX
AC AAZ072
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DE Human
XX
MY
DE Human
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DE Human
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MY
DO Synthe
OS Synthe
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PN WO993E
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ABL47224 standard; RNA; 17 BP

ABL47224;

ABL47224/c ID ABL472 XX AC ABL472 XX

RESULT 350

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The present invention relates to oligonucleotides that downregulate the expression of human GrD2-related with Insert Domain (GRID) gene. GRID is a T-cell co-stimulatory adaptor protein. The oligonucleotides are useful for modulating the expression of GRID, to treat conditions such as its sucygraft rejection and leukaemia. The oligonucleotides can also be administered in conjunction with other therapies such as radiation, chemotherapy and cyclosporin treatment. The present oligonucleotide was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                              Human, Grb2-related with Insert Domain; GRID; T-cell; co-stimulatory adaptor protein; tissue rejection; graft rejection; leukaemia; cytostatic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human, Grb2-related with Insert Domain; GRID; T-cell; co-stimulatory adaptor protein; tissue rejection; graft rejection; leukaemia; cytostatic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ellis JH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New nucleic acid(s) for regulating the Grb2-related with Insert D (GRID) gene comprises using antisense and enzymatic nucleic acid molecules such as hammerhead ribozymes.
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3.0%; Score 13.4; DB 1; Length 17;
Best Local Similarity 93.3%; Pred. No. 3.6e+02;
Matches 14; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mcswiggen JA, Hamblin PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human GRID NCH ribozyme substrate oligonucleotide #250.
Human GRID Amberzyme substrate oligonucleotide #124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17 BP; 3 A; 3 C; 10 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 4; Page 87; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               used to illustrate the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP
                                                                                                                                                                                                                                                                                          23-FEB-2001; 2001WO-US005957.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          146
                                                                                                                                                                                                                                                                                                                                          24~FEB-2000; 2000US-0184594P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-FEB-2001; 2001WO-US005957.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Jarvis T, Von Carlowitz I,
                                                                                                                                                                                                                                                                                                                                                                                        (RIBO-) RIBOZYME PHARM INC. (GLAX ) GLAXO GROUP LID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15 CTCGCCTGCCGCCT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABL46796 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        132 CTCGGCCTGCCGCCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-550088/61.
                                                                                                                                                                                               WO200162911-A2
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                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27-JUN-2003
                                                                                                                                                                                                                                             30-AUG-2001
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(DNAS-) DNA SCI INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAL44028;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 353
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAL44028
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                                                                                                                                                                   The present invention relates to oligonucleotides that downregulate the expression of human Grb2-related with Insert Domain (GRID) gene. GRID is a T-cell co-stimulatory adaptor protein. The oligonucleotides are useful for modulating the expression of GRID, to treat conditions such as tissue/graft rejection and leukaemia. The oligonucleotides can also be administered in conjunction with other therapies such as radiation, chemotherapy and cyclosporin treatment. The present oligonucleotide was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel polynucleotide which down regulates expression of Ets-related gene,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             neovascular glaucoma, myopic degeneration, arthritis, verruca vulgaris, angiofibroma of tuberous sclerosis, port-wine stain, wound healing; Sturge Weber syndrome, Kippel-Trenaunay-Weber syndrome, leukaemia, 88;
                                                                                                 New nucleic acid(s) for regulating the Grb2-related with Insert Domain (GRID) gene comprises using antisense and enzymatic nucleic acid molecules such as hammerhead ribozymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ophthalmological, antiarthritic, antipsoriatic, virucide, osteopathic, vulnerary, cancer, lymphoma; Ewing's sarcoma; melanoma; psoriasis; tumour angiogenesis; diabetic retinopathy; macular degeneration;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Osler-Weber-rendu syndrome, leukaemia; osteoporosis; DNAzyme; inozyme;
                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Randi AM;
                                                       Ellis JH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; hammerhead ribozyme; cytostatic; antitumour; antidiabetic;
                                                                                                                                                                                                                                                                                                                      ;
0
                                                                                                                                                                                                                                                                                              3.0%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 3.6e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Von Carlowitz I, Mcswiggen JA, Mclaughlin F,
                                                       Mcswiggen JA, Hamblin PA,
                                                                                                                                                                                                                                                                        Sequence 17 BP; 3 A; 4 C; 9 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human ERG DNAzyme target sequence Seq ID No 1481.
                                                                                                                                               Claim 4; Page 67; 108pp; English.
                                                                                                                                                                                                                                                  used to illustrate the invention
                                                                                                                                                                                                                                                                                                                                                                                                                         ABK18834 standard; RNA; 17 BP
24-FEB-2000; 2000US-0184594P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16-MAY-2001; 2001WO-US015866.
                                                                                                                                                                                                                                                                                                                                           132 CTCGGCCTGCCGCT 146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16-MAY-2000; 2000US-00572021
                                                       Von Carlowitz I,
                      (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                        CICGCCCIGCCGCCI 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (RIBO-) RIBOZYME PHARM INC (GLAX ) GLAXO GROUP LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                      14; Conservative
                               (GLAX ) GLAXO GROUP LTD
                                                                             WPI; 2001-550088/61.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-082995/11.
                                                                                                                                                                                                                                                                                                          Best Local Similarity
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                                                       Jarvis T,
                                                                                                                                                                                                                                                                                                                                                                  16
                                                                                                                                                                                                                                                                                                                                                                                                                                               ABK18834;
                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                    352
                                                                                                                                                                                                                                                                                                                      Matches
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conditions elected from cancer, lymphoma, Ewing's sarcoma, melanoma, conditions selected from cancer, lymphoma, Ewing's sarcoma, melanoma, conditions selected from cancer, lymphoma, Ewing's sarcoma, melanoma, cumour angiogeneration, archiritis, psoriasis, verruca vulgaris, angiofibroma of tuberous sclerosis, port-wine stains, Sturge correcting a patient having a condition associated with the level of ERG, treating a patient having a condition associated with the level of ERG, by contacting cells of the patient with (I) under conditions suitable for the treatment. Leukaemia or tumour conditions suitable for the treatment. Leukaemia or tumour conditions suitable for the treatment. Leukaemia or tumour conjunction with one or more of other therapies such as radiation or chemotherapy treatment. (I) is useful for reducing ERG activity in a conjunction such as Mg2+. (I) is useful for dispensed of contitions and diseases related to the expression of ERG, and as diagnostic tool to cation such as Mg2+. (I) is useful for specifically cation genes that share homology with ERG gene or ERG fusion genes. Constructing within diseased or ERG fusion genes. Constructing such as a cell. (I) is useful for specifically cation such as Mg2+. (I) is useful for specifically contacting genes that share homology with ERG gene or ERG fusion genes. Constructing such as mandament.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
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CYP5402A6; CYP2A6, chromosome 19; steroid metabolism;
drug detoxification; xenobiotic detoxification; procarcinogen activation;
inflammation; asthma; habitual smoking.
useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  enzymatic nucleic acid molecules which regulate expression of ERG, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Сарв
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human cytochrome P4502A6 (CYP4502A6) promoter - polymorphic region 3.
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/standard_name= "Single nucleotide polymorphism"
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                                                                                                                                                                                                                                          relates to a nucleic acid molecule (I) which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 17 BP; 1 A; 5 C; 6 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         related PCR primers of the invention
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replace(10, C)
                                                                                                                                           Claim 4; Page 93; 149pp; English.
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Best Local Similarity 93.3
Matches 14; Conservative
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ACA07669;
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                                                                                                                           The invention comprises the identification of genetic polymorphisms in the human cytochrome P4502A6 (CYP4502A6 or CYP2A6) gene. The human cytochrome P4502A6 gene is located on chromosome 19 and encodes an enzyme that plays a role in the metabolism of steroids, the detoxification of drugs and xenobiotics, and the activation of procarcinogens. The P4502A6 polymorphisms identified in the invention are useful for evaluating an individual's risk of developing asthma or an individual's propensity for cigarette consumption. The P4502A6 DNA sequences of the invention are useful for identifying individuals having a polymorphic genotype and to screen individuals for altered metabolism for cyclothrome P4502A6 substrates. The P4502A6 DNA sequences of the invention are also useful for identifying individuals who are at risk from inflammation, asthma, habitual smoking and diseases that result from environmental or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; ds; single nucleotide polymorphism; SNP; cytochrome; P4502A6; CYP4502A6; CYP2A6; chromosome 19; steroid metabolism; drug detoxification; xenobiotic detoxification; procarcinogen activation;
                                                                                                                                                                                                                                                                                  occupational exposures to dangerous substances. The present DNA sequence represents a polymorphic region of the promoter for the human cytochrome P4502A6 gene
                                                        New isolated polynucleotide, useful to screen individuals for asthma, inflammation and susceptibility to habitual smoking, comprises base variation from that of known human cytochrome P4502A6 sequence.
                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human cytochrome P4502A6 (CYP4502A6) promoter - polymorphic region 4.
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                                                                                                                                                                                                                                                                                                                                    Sequence 17 BP; 4 A; 5 C; 1 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           inflammation; asthma; habitual smoking.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                         Claim 35; Page 27; 48pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ВР
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                                    WPI; 2002-566448/60.
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                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
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            Guida M, Hall J;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                    14;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Key
variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAL44029;
                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 354
                                                                                                                                                                                                                                                                                                                                                                                    Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; G-cleaver; amberzyme; cancer; REL-A activity; breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; cospapageal cancer; stomach cancer; bladder cancer; pancreatic cancer; cervical cancer; nead and neck cancer; ovarian cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxel; docetaxel; cisplatin; methotrexate; cyclophosphamide; doxorubin; fluorouracil carboplatin; edatrexate; gemcitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatorid arthritis; restenosis; cohn's disease; obesity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammatory bowel disease; infection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               occupational exposures to dangerous substances. The present DNA sequence represents a polymorphic region of the promoter for the human cytochrome
                                                                  New isolated polynucleotide, useful to screen individuals for asthma, inflammation and susceptibility to habitual smoking, comprises base variation from that of known human cytochrome P4502A6 sequence.
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Pred. No. 3.6e+02;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17 BP; 4 A; 6 C; 1 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NFKB sub-unit modulating zinzyme substrate #68.
                                                                                                                                                                                                                                              Claim 1; Page 27; 48pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                92US-00987132.
94US-00245466.
94US-00291932.
96US-00777916.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 93.3
Matches 14; Conservative
WPI; 2002-566448/60.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US2002177568-A1.
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15-AUG-1994;
23-DEC-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       P4502A6 gene
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ABZ66567;
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                                                                                                                                                                                                                       The invention describes an enzymatic nucleic acid molecule (I) which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFRS), where (I) is an inozyme, zinzyme, G-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a patient having a condition associated with the level of REL-A. (I) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially MG-2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal, brain, oesophagel, stoomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug therapies such as monoclonal antibodies, REL-A-specific inhibitors or chemotherapy including paclitaxel, docetaxel, cisplatin, methotrexate, cyclophosphamide, doxorubin, fluorouzacil carboplatin, edatrexate, cyclophosphamide or radiation therapy. The enzymatic and antisense nucleic acid molecules are also useful for treating inflammatory disease such as theumatoid arthritis, restenosis, asthma. Crohn's disease, diabetes, obesity, autoimmune disease, lupus, multiple sclerosis, transplant/graft rejection, gene therapy applications, asthma. Central nervous system (CNS) and myocardial), glomerulonephritis, especial are allergic airway inflammation, inflammatory bowel disease or configuration and antisemed or configuration and ant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                              Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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enzymatic nucleic acid; H-Ra8; N-Ra8; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; 88.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3.0%; Score 13.4; DB 1; Length 17; 73.3%; Pred. No. 3.6e+02; tive 3; Mismatches 1; Indel8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 17 BP; 0 A; 9 C; 4 G; 0 T; 4 U; 0 Other;
                                                                 Stinchcomb DT, Mcswiggen J, Draper KG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human HIV DNA zyme substrate #13.
                                                                                                                                                                                            Claim 3; Page 38; 72pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABZ66556 standard; RNA; 17 BP
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 STINCHCOMB D T. MCSWIGGEN J.
                                                                                                WPI; 2003-340953/32.
                                 DRAPER K G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
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(STIN/)
                                 (DRAP/)
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ABZ66556
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The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule, that modulates expression of a nucleic acid molecule acid molecule, that modulates human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV acid molecules are useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ59889 - ABZ62216, ABZ64544 - ABZ65531, ABZ6520 - ABZ65224, ABZ65530 - ABZ65216, Human
                                                                                                                                                                                                                                                                                                                                            Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Seguence 17 BP; 3 A; 5 C; 6 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 123; Page 157; 185pp; English.
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06-JUN-2001; 2001US-0296249P.
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06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
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                                                                                                                (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ribozymes of the invention
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                                                                                                                                                                                            Mcswiggen J;
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(RIBO-) RIBOZYME PHARM INC.
                                                                                                    infection.
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                                                       Novel
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                                                                 The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-theumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are blacker, or pancreatic cancer, and HIV infection, and AIDS. The sequences bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ56858 represent substrate/target sequences for the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; hammerhead riboxyme; DNAzyme; inozyme; zinzyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative, disease state, HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
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  HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HCV minus strand DNAzyme substrate sequence #1361.
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                                                                                                                                                                                                                                                                                                                                                Seguence 17 BP; 3 A; 6 C; 6 G; 0 T; 2 U; 0 Other;
                                          Claim 123; Page 158; 185pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mcswiggen J,
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2001US-0296876P.
2001US-0335059P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ACD64058 standard; RNA; 17 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                 12; Conservative
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Roberts E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BLATT L.
MACEJAK D.
MCSWIGGEN J.
MORRISSEY D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PAVCO P.
LEE P.
DRAPER K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ROBERTS E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200281494-A1.
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08-JUN-2001; 2
08-JUN-2001; 2
24-OCT-2001; 2
05-DEC-2001; 2
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Draper K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ACD64058;
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(BLAT/)
(MACE/)
(MCSW/)
(MORR/)
(PAVC/)
(LEEP/)
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(ROBE/)
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ACD64058/C
ID ACD64058/C
XX ACD640;
XX ACD640;
XX ACD640;
XX ACD 30-SEP
XX ACD640;
XX ACCCT
XX ACC
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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes. DNAzymes, concerning are nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptase and/or HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV DNA. The nucleic acids may be used to modulate the expression of HBV CO DNA. The nucleic acids may be used to modulate the expression of HBV CO DNA. The modulate the expression Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and compounds of disease states related to HBV and HCV infection, replication and gene concession such as cirrhosis, liver failure, and hepatocallular carcinoma. The present sequence represents a substrate for one of the HCV DNAzyme or minus strand DNAzyme sequences disclosed in the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                              Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nucleic acid molecule, Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; hammerhead ribozyme; DNAzyme; inozyme; zinzyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptemer; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3.0%; Scc. 93.3%; Pred. No. 5...
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                                                                                                                                                                                                                     Claim 1; Page 299; 387pp; English.
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08-JUN-2001, 2001US-0087478.
08-JUN-2001, 2001US-0296876P.
24-OCT-2001, 2001US-0335659P.
05-DEC-2001, 2001US-0337055P.
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WPI; 2003-229207/22.
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Best Local Similarity
Matches 14; Conserv
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17-SEP-2001; 2001FR-00011980.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 361
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ð
                                                                                                                                                                                                                                                                                                                                      The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HVV) or Hepatitis B virus (HVV) or Apatitis B virus (HVV) or an oncleic acide and an entisense and enzymatic nucleic acids such as hammerhead ribozymes. DNAzymes, inozymes, zinzymes, amberzymes, and G-cleaver ribozymes. Also disclosed are and eacy molecules and aptamers that bind to HBV reverse transcriptase primar sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV DNA. The nucleic acids may be used to modulate the expression of HBV compounds and/or potential theraples directed against HBV, and compounds compounds and/or potential theraples directed against HBV, and compounds compounds of the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene compounds and the compounds and HCV infection, replication and gene configuration and gene acids may be accounted and the procession and specific acids and repatication and gene acids may be accounted and seasons and acids acids and acids and acids acids and acids acids and acids and acids and acids acids acids and acids acids acids acids and acids acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      tumour suppression; tumour reversion; apoptosis; virus resistance; yctostatic; virucide; neuroprotective; nootropic; neuroleptic; probe; primer; PCR; gene chip; antisens; viral disease; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; ds; human.
                                                                                                                                                                                                                                        Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                      <u>ب</u>
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                                                                                                                                                      Lee
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3.0%; Score 13.4; DB 1; Length 17; 80.0%; Pred. No. 3.6e+02; tive 2; Mismatches 1; Indel8
                                                                                                                                                      Pavco P,
                                                                                                                                                      Mcswiggen J, Morrissey D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Seguence 17 BP; 1 A; 7 C; 7 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                            Claim 1; Page 250; 387pp; English.
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                                                                                                                                                     Macejak D,
Roberts E;
BLATT L.
MACEJAK D.
MCSWIGGEN J.
MORRISSEY D.
                                                                                                                                                                                                        WPI; 2003-229207/22
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                                                                 PAVCO P.
LEE P.
DRAPER K.
                                                                                                                     ROBERTS E.
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                                                                                                                                                                                                                                                                            infection.
                                                                                                                                                                          Draper K,
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                                                                                                                                                   Blatt L,
                               (MCSW/) N
(MORR/) N
(PAVC/) I
(LEEP/) I
(DRAP/) I
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This invention relates to novel isolated nucleic acid sequences involved in the phenomena of tumour suppression, tumour reversion, apoptosis and/or resistance to viruses. The invention may be useful for the development of compounds with a cytostatic, virucide, neuroprotective, notropic or neuroleptic activity. The DNA sequences may be useful as probes and primers for detecting, indentifying, quantifying and/or complication for example as one component of a gene chip, in viro as antisense reagents and for production of recombinant of polypeptides. The invention may therefore be useful for preparation of polypeptides. The invention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia. The present sequence is that of a nucleic acid sequence of the invention.

Once: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                          New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.
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NCH ribozyme, G-cleaver ribozyme, Zinzyme, DNAzyme, amberzyme, Inozyme,
hairpin ribozyme, tissue rejection, graft rejection, leukaemia.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
3.0%; Score 13.4; DB 1; Length 17;
Best Local Similarity 93.3%; Pred. No. 3.6e+02;
Matches 14; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                   Disclosure; SEQ ID NO 1134; 30pp; French
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(MOLE-) MOLECULAR ENGINES LAB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  97 TGTTTTTCTCGCTGA 111
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MCSWIGGEN J.
HAMBLIN P A.
ELLIS J H.
                                                                    felerman A, Amson R,
                                                                                                                                         WPI; 2003-313354/30.
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(MCSW/)
(HAMB/)
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US09436060A-16.rng.sl

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The invention relates to a nucleic acid molecule that down-regulates expression of Grb2-related with insert domain (GRID) gene, e.g. a harmerhead ribozyme, NCH ribozyme, G-cleaver ribozyme, Zinzyme, DNAzyme, anderhead ribozyme, NCH ribozyme, Also include are a mammalian cell including the novel nucleic acid molecule, reducing GRID activity in a cell by contacting the cell with the novel nucleic acid molecule, treating a patient having a condition associated with the level of GRID (e.g. tissue/graft rejection or leukaemia) by contacting the cell with the novel nucleic acid molecule, cleaving RNA of a GRID gene by contacting the cell with the novel nucleic acid molecule, an expression vector comprising a nucleic acid sequences (encoding at least the novel nucleic acid molecule in a manner that allows its expression), a manner that allows its expression), a manner that cell including the expression vector and an enzymatic nucleic acid molecule that cleaves RNA derived from a GRID gene. The nucleic acid molecule that releaves a condition associated with the level of GRID, e.g. tissue/graft rejection and leukaemia. The present sequence is a target region for the enzymatic nucleic acids of the invention.
                                                New nucleic acid molecule that down-regulates expression of Grb2-related with insert domain (GRID) gene, useful for treating a condition associated with the level of GRID, e.g. tissue/graft rejection and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    enzymatic nucleic acid; RNA cleavage; hepatitis C virus; HCV; infection; type I interferon; DNAzyme.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17 BP; 3 A; 3 C; 10 G; 0 T; 1 U; 0 Other;
                                                                                                                                    Claim 4; SEQ ID NO 857; 74pp; English.
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MCSWIGGEN J.
ROBERTS E.
PAVCO P A.
MACEJACK D.
                  WPI; 2003-829646/77
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                                                                                                     leukemia.
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(ROBE/)
(PAVC/)
(MACE/)
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Matches
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Enzymatic nucleic acid molecules which specifically cleave RNA derived

Macejack D;

Pavco PA,

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from hepatitis C virus (HCV), useful for the treatment of HCV infections, especially in combination with type I interferon therapy.
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                                                                         The invention relates to an enzymatic nucleic acid molecule which specifically cleaves RNA derived from hepatitis C virus (HCV), in which the binding arms of the enzymatic nucleic acid molecule comprises sequences complementary to any of the defined substrate sequences given in the specification. The nucleic acid molecule may be administered for the treatment of HCV infections, especially in combination with type I interferons. The present sequence represents a HCV DNAzyme substrate
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/mod_base= OTHER
/note= "Sugar-phosphate backbone has been replaced by
peptide backbone"
                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                 Sequence 17 BP; 2 A; 7 C; 6 G; 0 T; 2 U; 0 Other;
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                                             Claim 1; SEQ ID NO 3638; 198pp; English.
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(first entry)
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Les 14; Conserv
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modified_base
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Conservative
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12-MAY-1998
13;
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                                    13
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AAZ08815/c
Matches
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mammalian telomerase and as inhibitors of telomerase activity, especially
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/*tag= a
/mod_bage= OTHER
/note= "Sugar-phosphate backbone has been replaced by opptide backbone"
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                           Revised record issued on 21-OCT-2004 : Correction to feature table key
                                                                                      Gaps
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                                                                 Score 13; DB 1; Length 13;
Pred. No. 2.8e+02;
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                                                                                     0; Indels
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                                               Sequence 13 BP; 5 A; 1 C; 4 G; 3 T; 0 U; 0 Other
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                                                          2.9%; Scc...
100.0%; Pre
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                                                                                                         42 TIGICIAACCCIA 54
                                                                                      Conservative
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Best Local Similarity
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12-MAY-1998
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Best Local
                                                                                                                                                                                                                                                                   Peptide
                                                                                                                                                        RESULT 364
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New peptide nucleic acids hybridising to mammalian telomerase RNA - used to inhibit telomerase, for treating tumours and other proliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /nore= "sugar-phosphate backbone has been replaced by a peptide backbone"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Norton JC;
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/mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            96US-00630019.
                                                                                                                                                                                                  AAT89225 standard; DNA; 13
                                                                                                                                                                                                                                                                                                               (revised)
(first entry)
44 GTCTAACCCTAAC 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        46 CTAACCCTAACTG 58
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Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CTAACCCTAACTG 1
                                            GTCTAACCCTAAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Wright WE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1997-512647/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Key
modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GERO-) GERON
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US09436060A-16.rng.sl

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The present sequence represents a peptide nucleic acid molecule which hybridises to the mRNA component of mammalian telomerase, and inhibits telomerase activity. Telomerase is a ribonucleoprotein enzyme that synthesizes one strand of the telomeric DNA, using as a template an ill conclectes now strand of the telomeric DNA, using as a template an ill conclectide sequence contained within the RNA component of the enzyme. The invention relates to PNA molecules having a sequence of no more than 25 concers, which include the sequence of TTAGG. The uncharged nature of the PNA increases the melting temperature of associating strands, increases the melting temperature of associating strands, increases the rate of association with targeted nucleic acids, and affords greater resistance of degradation by proteases or nucleases. The corrects, neoplasia, hyperplasia, neurodegenerative diseases. The affords greater resistance of degradation by proteases conditions such as cancers, neoplasia, hyperplasia, neurodegenerative diseases. The diseases haracterized by abnormal telomere metabolism or telomerase diseases haracterized by abnormal telomere metabolism or telomerase diseases haracterized by abnormal telomer metabolism or telomerase cartivity, in combination with antineoplastic and other cytotoxic or cytostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization probes to detect the RNA component of a mammalian telomerase and as inhibitors of telomerase activity. The method of the present invention to be detected with increase and as inhibitors of telomerase activity. The method of the present invention can be used to detect an immortal or mechods of the present invention can be used to detect an immortal or mechods of the present invention can be used to detect an immortal or cell expresses telomerase activity and its RNA component
                                                                                                                                                                                                                                                                                                                                          New peptide nucleic acid (PNA) compounds that inhibit telomerase activity in mammalian cells is useful as probes to detect the RNA component of a mammalian telomerase.
via glycine amino N through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                            Corey DR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.9%; Score 13; DB 1; Length 13;
LOO.0%; Pred. No. 2.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PNA sequence #52 used to inhibit telomerase activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 13 BP; 3 A; 1 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                            Wright WE, Piatyszek MA, Shay JW, Norton JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Prec. ... 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                   6; Col 71; 45pp; English.
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                                                                                                                                                                         96US-00630019
                                                                                                                               97US-00838545
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                 (TEXA ) UNIV TEXAS SYSTEM
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Matches 13; Conservative
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                                                                                                                                                                                                                                                                                                      WPI; 2000-292432/25.
                                                                                                                               09-APR-1997;
                                                                                                                                                                         09-APR-1996;
                                              US6046307-A
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                                                                                      04-APR-2000
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AAA37594/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes an antitumour augmenting agent comprising a hybridisable oligonucleotide, as an effective component, contained at the template region of RNA site contained in telomerase. Also described
                                                                                                                                                                       Human; RERF-LC-A1; hybridisation; antitumour augmenting drug; telomerase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Peptide nucleic acid; PNA; telomerase; ribonucleoprotein enzyme; cancer; inhibitor; neoplasia; neurodegenerative disease; aging; hyperplasia; AIDS; HIV; fungal infection; forensic identification; detect; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            is an antitumour composition comprising the antitumour augmenting agent and an antitumour agent as effective components. The present sequence represents a hybridisable oligonucleotide for human RERF-LC-A1, used in the exemplification of the present invention. When the antitumour composition is administrated into a mammal, especially human, a sufficient antitumour effect is exhibited to cancer cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /*tag= a //rag= a mucleic acid molecule, where N-(2-aminoethyl)glycine units are linked to nucleotide bases
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New composition - for anti-tumor augmenting agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Seguence 13 BP; 5 A; 1 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                               Human RERF-LC-Al hybridisable oligonucleotide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Page 6; 8pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (KURE ) KUREHA CHEM IND CO L'TD
    ВР
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    AAZ08815 standard; DNA; 13
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                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     paternity testing; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1999-522625/44
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                                                                                                                                                                                                                                                                Homo sapiens.
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misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                           10-FEB-1998;
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                                                                                      01-NOV-1999
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                                                                                                                                                                                                                                                                                                                                                24-AUG-1999.
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                                                                                                                                                                                                    cancer; 88
                                                                                                                                                                                                                                          Synthetic
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                                              AAZ08815
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Gaps

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Peptide nucleic acid, DNA; telomerase, ribonucleoprotein enzyme; cance
inhibitor; neoplasia; neurodegenerative disease; aging; hyperplasia;
AIDS; HIV; fungal infection; forensic identification; detect; tumour;
                                  Location/Qualifiers
                paternity testing; ss.
                                                                                                                     WPI; 2000-292432/25.
                                     misc_feature
                                                                                  09-APR-1997;
                                                                                           09-APR-1996;
                                                                 US6046307-A.
                                                                         04-APR-2000.
                         Synthetic
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nucleotice sequence contained within the KNA component of the anzyme. The invention relates to PNA molecules having a sequence of no more than 25 bases, which include the sequence GTTAGG. The uncharged nature of the PNA backbone increases the melting temperature of associating strands, increases the rate of association that targeted nucleic acids, and affords greater resistance of degradation by proteases or nucleases. The increases the rate of degradation by proteases or nucleases. The cherapeutic PNAs may be used for treating disease conditions such as cancers neoplasia, hyperplasia, neurodegenerative diseases, aging, human immondeficiency virus (HIV) infection/AIDS (acquired immunodeficiency virus (HIV) infection/AIDS are used for molecular diagnostics, labelled PNAs are used as hybridization or probes to detect or quantitate polymucleotides having a human telomerase and somitividuals. e.g. paternity testing, based on HTR sequence. PNA probes are also used for forensic identification of fragment length polymorphism (RPLP) pattern. PNAs are also useful as inhibitors of telomerase activity. The method of the present invention allows cancerous conditions to be detected with increased confidence and allows cancerous conditions to be detected with increased confidence and allows cancerous conditions to be detected with increased confidence and allows cancerous conditions to be detected with increased confidence and based on pathological characteristics. The diagnostic an immortal or methods of the present invented according any origin, 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      hybridiess to the mRNA component of mammalian telomerase, and inhibits telomerase activity. Telomerase is a ribonucleoprotein enzyme that synthesizes one strand of the telomeric DNA, using as a template an 11 nucleotide sequence contained within the RNA component of the enzyme. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New peptide nucleic acid (PNA) compounds that inhibit telomerase activity in mammalian cells is useful as probes to detect the RNA component of a mammalian telomerase.
/*tag= a //*hote= "Peptide nucleic acid molecule, where N-(2-aminoethyl)glycine units are linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  present sequence represents a peptide nucleic acid molecule which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Corey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 13 BP; 7 A; 1 C; 3 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Wright WE, Piatyszek MA, Shay JW, Norton JC,
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The present sequence represents a peptide nucleic acid molecule which hybridises to the mRNA component of mammalian telomerase, and inhibits ctelomerase activity. Telomerase is a ribonucleoprotein enzyme that extended the telomeric DNA, using as a template an 11 sourceotide sequence contained within the RNA component of the enzyme. The synthesizes one strand of the telomeric DNA, using as a template an 11 nucleotide sequence contained within the RNA component of the enzyme. The invention relates to PNA molecules having a sequence of no more than 25 backbone increases the melting temperature of associating strands, increases the rate of association with targeted nucleic acids, and affords greater resistance of degradation by proteases or nucleases. The charapeutic PNAs may be used for treating disease conditions such as cancers, neoplasia, hyperplasia, neurodegenerative diseases. The characterized by abnormal telomere metabolism or telomerase communication with antineoplastic and other cytocoxic or sectivity, in combination with antineoplastic and other cytocoxic or cytostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization or cytostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization of individuals, e.g. paternity testing, based on hTR gene restriction of individuals, e.g. paternity testing, based on hTR gene restriction fragment length polymorphism (RFLP) pattern. PNAs are also useful as probes to detect the RNA component of a mammalian telomerase and as probes to detect the RNA component of a mammalian telomerase and as inhibitors of telomerase activity. The method of the present invention allows cancerous conditions to be detected with increased confidence and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New peptide nucleic acid (PNA) compounds that inhibit telomerase activity
                                                                                                                                                                                                                                                                                                              cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   in mammalian cells is useful as probes to detect the RNA component of a mammalian telomerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note= "Peptide nucleic acid molecule, where N-(2-aminoethyl)glycine units are linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                         Peptide nucleic acid; PNA; telomerase; ribonucleoprotein enzyme; cance
inhibitor; neoplasia; neurodegenerative disease; aging; hyperplasia;
AIDS; HIV; fungal infection; forensic identification; detect; tumour;
paternity testing; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Corey DR;
                                                                                                                                                                                                                                                               PNA sequence #56 used to inhibit telomerase activity.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
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                                                                                                                                 AAA37598 standard; DNA; 13 BP.
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                  TTTTTGTCTAACC
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WPI; 2000-292432/25.
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                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                       13
                                                                                                                                                                          AAA37598;
                                                                                                       AAA375987
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Query Match 2.9 Best Local Similarity 100. Matches 13; Conservative

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Gaps

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2.9%; Score 13; DB 1; Length 13; 100.0%; Pred. No. 2.8e+02; ive 0; Mismatches 0; Indels

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New peptide nucleic acid (PNA) compounds that inhibit telomerase activity in mammalian cells is useful as probes to detect the RNA component of a mammalian telomerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Peptide nucleic acid, PNA, telomerase, ribonucleoprotein enzyme; cancer, inhibitor; neoplasia; neurodegenerative disease; aging; hyperplasia; AIDS, HIV; fungal infection; forensic identification; detect; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /,oct= "Peptide nucleic acid molecule, where N-(2-aminoethyl)glycine units are linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
possibly at an earlier stage, before cells are detected as cancerous based on pathological characteristics. The diagnostic and prognostic methods of the present invention can be used to detect an immortal or neoplastic cell or tumour tissue or cancer of any origin, provided the cell expresses telomerase activity and its RNA component.
                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Corey DR;
                                                                                                                                     Length 13;
                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                      PNA sequence #51 used to inhibit telomerase activity.
                                                                                                  Sequence 13 BP; 1 A; 6 C; 2 G; 4 T; 0 U; 0 Other;
                                                                                                                                     Score 13; DB 1; Le
Pred. No. 2.8e+02;
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                                                                                                                                              100.0%; Preα. ...
'''e 0; Mismatches
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                                                                                                                    55 ACTGAGAAGGGCG 67
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                                                                                                                                                                                                                                      ACTGAGAAGGGCG 1
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immunodeficiency virus (HIV) infection/AIDS (acquired immunodeficiency
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"Phosphorothioate internucleotide linkages"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 13; DB 1; Length 13;
Pred. No. 2.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         38 TITITIGICIAAC 50
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Les 13; Conservative
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/note=
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Matches
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controlled sequence contained within the RNA component of the enzyme the nucleotide sequence contained within the RNA component of the enzyme. The nucleotide sequence contained within the RNA component of the enzyme. The nucleotide sequence contained having a sequence of no more than 25 bases, which include the sequence GTTAGG. The uncharged nature of the PNA backbone increases the melting temperature of associating strands, controlled the sequence of degradation with targeted nucleic acids, and affords greater resistance of degradation by proteases or nucleases. The therapeutic PNAs may be used for treating disease conditions such as cancers, neoplasia, hyperplasia, neurodegenerative diseases. adjing, human immunodeficiency virus (HIV) infection/AIDS (acquired immunodeficiency syndrome) and associated pathologies, fungal infections, and other cycostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization cycostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization of individuals, e.g. paternity testing, based on hTR gene restriction of individuals, e.g. paternity testing, based on hTR gene restriction contained the RNA component of a mammalian telomerase and as thibbitors of telomerase activity. The method of the present invention allows cancerous conditions to be detected with increased confidence and
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control sequence alongside a peptide nucleic acid molecule which hybridises to the mRNA component of mammalian telomerase, and inhibits telomerase activity. Telomerase is a ribonucleoprotein enzyme that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      possibly at an earlier stage, before cells are detected as cancerous based on pathological characteristics. The diagnostic and prognostic methods of the present invention can be used to detect an immortal or neoplastic cell or tumour tissue or cancer of any origin, provided the cell expresses telomerase activity and its RNA component
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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AAA37597 standard; DNA; 13 BP.
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Best Local Similarity
Matches 13; Conserv
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The present sequence represents a peptide nucleic acid molecule which hybridises to the mRNA component of mammalian telomerase, and inhibits call hybridises to the mRNA component of mammalian telomerase, and inhibits call hybridises to the mRNA component of mammalian telomerase, and inhibits of synthesizes one strand of the telomeric DNA, using as a template an 11 nucleotide sequence contained within the RNA component of the enzyme. The invention relates to PNA molecules having a sequence of no moore than 25 bases, which include the sequence drawing a sequence of no moore than 25 chases, which include the sequence drawing a sequence of the PNA carefords increases the melting temperature of associating strands, increases the rate of association with targeted nucleic acids, and affords greater resistance of degradation by proteases or nucleases. The charapeutic PNAs may be used for treating disease conditions such as cancers, neoplasia, hyperplasia, neurodegenerative diseases or nucleases. The charapeutic PNAs may be used for memore and other manned ficiency virus (HIV) infection/AIDS (acquired immunodeficiency virus (HIV) antineoplastic and other cytcoxic or cytostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization of individuals, e.g. paternity testing, based on HTR gene restriction of individuals, e.g. paternity testing, based on HTR gene restriction of individuals, and component of a mammalian telomerase and as thinbitors of telomerase activity. The method of the present invention allows cancerous conditions to be detected with increased conditions and allows cancerous conditions to be detected with increased conditions.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          possibly at an earlier stage, before cells are detected as cancerous based on pathological characteristics. The diagnostic and prognostic methods of the present invention can be used to detect an immortal or neoplastic cell or tumour tissue or cancer of any origin, provided the
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                                                                          Norton JC,
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100.0%; Pred. No. 2.8e+02;
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                                                                       Shay JW,
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                                                                          Piatyszek MA,
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                         (TEXA ) UNIV TEXAS SYSTEM
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Best Local Similarity 100.
Matches 13; Conservative
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                                                                                                                        WPI; 2000-292432/25.
                                                                          Wright WE,
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The present sequence represents a peptide nucleic acid molecule which hybridises to the mRNA component of mammalian telomerase, and inhibits to hybridises to the mRNA component of mammalian telomerase, and inhibits telomerase activity. Telomerase is a ribonucleoprotein enzyme that synthesizes one strand of the telomeric DNA, using as a template an 11 nucleotide sequence contained within the RNA component of the enzyme. The include the sequence of TTAGG. The uncharged nature of the PNA bases, which include the sequence of associating strands, increases the rate of association with targeted nucleic acids, and affords greater resistance of degradation by proteases or nucleases. The therapeutic PNAs may be used for treating disease conditions such as cancers, neoplasia, hyperplasia, neurodegenerative diseases. The therapeutic PNAs may be used for treating disease conditions such as cancers, neoplasia, hyperplasia, neurodegenerative diseases. The diseases haracterized by abnormal telomer and other cytotoxic or syndrome) and associated pathologies, fungal infections, and other diseases characterized by abnormal telomer and other cytotoxic or cytostatic agents, antifungal agents, and other nucleotides. PNAs may be activity, in combination with antinophalatic and other cytotoxic or cytostatic agents, antifungal agents, and other nucleotides. PNAs may be used for molecular diagnostics, labelled PNAs are used as hybridization cytostatic agents. The diagnostic and propers and as coopes to detect the RNA component of a mammalian telomerase and as probes to detect the RNA component of a mammalian telomerase and as inhibitors of telomerase activity. The method of the present invention can be used to detect as cancerous conditions to be detected with increased confidence and companient or allows cancerous conditions to be detected with increased confidence and companient or bused to detect an immortal or methods of the present invention can be used to detect an immortal or methods of the present invention can be used to detec
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mammalian cells is useful as probes to detect the RNA component of a
                      /*tag= a // hote= neptide nucleic acid molecule, where N-(2-aminoethyl)glycine units are linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                                                                                                   Corey DR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 13 BP; 3 A; 1 C; 5 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                 Shay JW, Norton JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 6; Col 71; 45pp; English.
                                                                                                                                                                                                                                     97US-00838545.
                                                                                                                                                                                                                                                                               96US-00630019.
                                                                                                                                                                                                                                                                                                                                                                                   Piatyszek MA,
                                                                                                                                                                                                                                                                                                                                 (TEXA ) UNIV TEXAS SYSTEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               mammalian telomerase.
                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-292432/25.
misc_feature
                                                                                                                                                                                                                                     09-APR-1997;
                                                                                                                                                                                                                                                                                    09-APR-1996;
                                                                                                                                         JS6046307-A
                                                                                                                                                                                        04-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                   Wright WE,
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                                   Gaps
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2.9%; Score 13; DB 1; Length 13;
100.0%; Pred. No. 2.8e+02;
ive 0; Mismatches 0; Indels
                                                                 44 GTCTAACCCTAAC 56
                     Local Similarity 100.
nes 13; Conservative
       Query Match
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AAA37547 standard; DNA; 13 BP.
GICTAACCCTAAC 1
       13
                                                                          AAA37547;
                                    RESULT 374
                                             AAA37547/c
ID AAA37
XX
AC AAA37
      g
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New peptide nucleic acid (PNA) compounds that inhibit telomerase activity in mammalian cells is useful as probes to detect the RNA component of a
                                       Peptide nucleic acid; PNA; telomerase; ribonucleoprotein enzyme; cancer;
                                                                                                                   /*tag= a //note= "Peptide nucleic acid molecule, where N-(2-aminoethyl)glycine units are linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
                                                inhibitor, neoplasia, neurodegenerative disease, aging, hyperplasia, AIDS, HIV, fungal infection, forensic identification, detect, tumour,
                                                                                                                                                                                                                                                  Corey DR;
                       PNA sequence #4 used to inhibit telomerase activity.
                                                                                                                                                                                                                                                  Shay JW, Norton JC,
                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                 96US-00630019.
                                                                                                                                                                                                 97US-00838545
                                                                                                                                                                                                                                                   Piatyszek MA,
       (first entry)
                                                                                                                                                                                                                                  (TEXA ) UNIV TEXAS SYSTEM.
                                                                   paternity testing; ss.
                                                                                                                                                                                                                                                                    WPI; 2000-292432/25.
                                                                                                                                                                                                 09-APR-1997;
                                                                                                                                                                                                                  09-APR-1996;
                                                                                                            misc_feature
       15-AUG-2000
                                                                                                                                                               JS6046307-A
                                                                                                                                                                                 04-APR-2000
                                                                                                                                                                                                                                                   Wright WE,
                                                                                   Synthetic.
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mammalian telomerase.

Example 1; Col 28; 45pp; English.

The present sequence represents a peptide nucleic acid molecule which hybridises to the mRNA component of mammalian telomerase, and inhibits calomerase activity. Telomerase is a ribonucleoprotein enzyme that exhibite stands of the telomeric DNA, using as a template an 11 invention relates to PNA molecules having a sequence of no more than 15 bases, which include the sequence drawing a sequence of no more than 25 bases, which include the sequence of TARGG. The uncharged nature of the PNA backbone increases the melting temperature of associating strands, increases the rate of association with targeted nucleid acids, and affords greater resistance of degradation by proteases or nucleases. The therapeutic PNAs may be used for treating disease conditions such as cancers, neoplasia, hyperplasia, neurodegenerative diseases. Aging, human immunodeficiency virus (HIV) infection/AIDS (acquired immunodeficiency probes to detect or quantitate, polymucleotides having a human telomerase RNA (HTR) sequence. PNA probes are also used for forensic identification of individuals, e.g. paternity testing, based on hTR gene restriction fragment length polymorphism (RFLP) pattern. PNAs are also useful as probes to detect the RNA component of a mammalian telomerase and as inhibitors of telomerase activity. The method of the present invention allows cancerous conditions to be detected with increased confidence and possibly at an earlier stage, before cells are detected as cancerous based on pathological characteristics. The diagnostic and prognostic methods of the present invention can be used to detect an immortal or neoplastic cell or tumour tissue or cancer of any origin, provided the cell expresses telomerase activity and its RNA component

Sequence 13 BP; 5 A; 1 C; 4 G; 3 T; 0 U; 0 Other;

DB 1; Length 13; 2.9%; Score 13;

Query Match

US09436060A-16.rng.sl

diseases characterised by abnormal telomere metabolism or telomerase

Page 191

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The present invention relates to peptide nucleic acids (PNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in mammalian cells by hybridising to the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polynucleotide having the human telomerase RNA component (hTR) sequence, as well as in forensic identification of individuals, such as peternity testing or identification of criminal suspects or unknown descendants based on the hTR gene RFLP pattern. The suspects or unknown descendants based on the hTR gene RFLP pattern. The pwphoproliferative diseases, and component diseases, or neurodegenerative diseases, and combination with other pharmaceuticals (such as antimoplastic or cytostatic agents) can be used for treating neoplasia, hymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases, automination with other pharmaceuticals (such as hyperplasia, human immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other
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/note= "This sequence is a peptide nucleic acid, i.e. it
contains a polyamide backbone instead of a deoxyribose
backbone"
                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or
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                                                                                                                                                                                                                                                                                        PNA 12 inhibiting human and mammalian telomerase activity.
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                  Indels
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 Pred. No. 2.8e+02;
                Mismatches
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               0 ;
                                                                                                                                                                                AAS15468 standard; DNA; 13 BP.
100.08;
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                                                    54
                  13; Conservative
                                                                               TTGTCTAACCCTA 1
                                                    42 TIGICIAACCCIA
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 Best Local Similarity
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Synthetic.
                                                                                                                                                                                                                  AAS15468;
                Matches
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     activity. The present sequence represents one of the PNA sequences of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to peptide nucleic acids (PNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in mammalian cells by hybridising to the RNA component of mammalian telomerase. The PNAs are useful as probes to detect the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polynucleotide having the human telomerase RNA component (hTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal
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/*tag=
/note= "This sequence is a peptide nucleic acid, 1.e. i
contains a polyamide backbone instead of a deoxyribose
backbone"
                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or
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                                                                                                                                                                                                                                                                                                                                                                     PNA 5/XII inhibiting human and mammalian telomerase activity.
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                                                                                           Length 13;
                                                                                                                         0; Indels
                                                             Sequence 13 BP; 2 A; 5 C; 1 G; 5 T; 0 U; 0 Other;
                                                                                                          2.8e+02;
                                                                                             DB 1;
                                                                                        2.9%; Score 13; DB 100.0%; Pred. No. 2.8 iive 0; Mismatches
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                                                                                                                                                                                                                                                                       ВР.
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                                                                                                                                                                                                                                                                       AAS15426 standard; DNA; 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       neurodegenerative diseases.
                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                                                                       53 TAACTGAGAAGGG 65
                                                                                                          Local Similarity 100.
                                                                                                                                                                                       TAACTGAGAAGGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Shay JW, Wright WE,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               modified base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            38-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                       AAS15426;
                                                                                           Query Match
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                                                                                                                           Matches
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Dynamic of the page of treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. The PMAs in combination with other pharmaceuticals (such as antineoplastic or cytostatic agents) can be used for treating neoplasia, hyperplasta, human immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomere metabolism or telomerase activity. The present sequence represents one of the PNA sequences of the invention. Note: The present sequence represents SED ID No 4 in the SEQ ID listing and column 4. However, table 1 in column 29 shows three sequences with SEQ ID No 4. The first SEQ ID No 4 (PNA #6, AASI5433) appears to be identical to SEQ ID No 1 in the claims and the SEQ ID isting. The second SEQ ID No 4 in table 1 appears to be identical to SEQ ID in the SEQ ID it the present sequence and appears to be identical to SEQ ID viet the present sequence and appears to be identical to SEQ ID viet the present sequence and appears to be identical to SEQ ID viet is the present sequence and appears to be identical to SEQ ID viet is the present sequence.
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    .13
    /*tag= a
    /*note="This sequence is a peptide nucleic acid, i.e. it contains a polyamide backbone instead of a deoxyribose backbone"

or unknown descendants based on the hTR gene RFLP pattern. The
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mammalian, peptide nucleic acid; probe; forensic; paternity testing; human telomerase RNA component; hTR gene RFLP pattern; cancer; inflammation; lymphoproliferative disease; autoimmune disease; neurodegenerative disease; neoplasia; hyperplasia; HIV; AIDS; human immunodeficiency virus; acquired immunodeficiency syndrome; telomere metabolism; mutant; cytostatic; anti-inflammatory; immunosuppressive; polyamide backbone; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 13; DB 1; Length 13;
Pred. No. 2.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 13 BP; 5 A; 1 C; 4 G; 3 T; 0 U; 0 Other;
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Best Local Similarity
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modified base
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09-APR-1997;
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New peptide nucleic acids that hybridizes to the RNA component of

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The present invention relates to peptide nucleic acids (PNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in nammalian cells by hybridising to the RNA component of mammalian telomerase and as inhibit or of telomerase activity, or to detect and/or quantitate polynucleotide having the human telomerase RNA component (HTR) sequence, as well as in forensic identification of component (HTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the hTR gene RFLP pattern. The suspects or unknown descendants based on the hTR gene RFLP pattern. The suspects or unknown descendants or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmaceuticals (such as antineoplastic or cyclostatic agents) can be used for treating neoplasia, hyperplasia, human immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomera metabolism or telomerase.
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                                                                                                                                                                                                                                                                                                                                                                                                                                     activity. The present sequence represents one of the PNA sequences of the
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/note= "This sequence is a peptide nucleic acid, i.e. it
contains a polyamide backbone instead of a deoxyribose
backbone"
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                inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases.
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mammalian telomerase, useful for treating or preventing cancer,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.9%; Score 13; DB 1; Length 13; 100.0%; Pred. No. 2.8e+02; cive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 13 BP; 3 A; 1 C; 5 G; 4 T; 0 U; 0 Other;
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                                                                            Claim 7; Col 73; 46pp; English.
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Best Local Similarity 100.
Matches 13, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-APR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              nvention
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                                                                                                                                                                  The present invention relates to peptide nucleic acids (PNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in mammalian cells by hybridising to the RNA component of mammalian telomerase and as inhibitors of telomerase activity or to detect and/or quantitate polynucleotide having the human telomerase RNA component (HTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the hTR gene RFLP pattern. The suspects or unknown descendants based on the hTR gene RFLP pattern. The suspects or unknown descendants or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmaceuticals (such as antineoplastic or cytostatic agents) can be used for treating neoplasia, hyperplasia, human immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other calesases characterised by abnormal telomere metabolism or telomerase activity. The present sequence represents one of the PNA sequences of the
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/note= "This sequence is a peptide nucleic acid, i.e. it
contains a polyamide backbone instead of a deoxyribose
backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mammalian; peptide nucleic acid; probe; forensic; paternity testing; human telomerase RNA component; hTR gene RPLP pattern; cancer; inflammation; lymphoproliferative disease; autoimmune disease; neurodegenerative disease; neoplasia; hyperplasia; HIV; AIDS; human immunodeficiency virus; acquired immunodeficiency syndrome;
                                                                          New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or
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                          Corey DR, Norton JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.9%; Score 13; DB 1; Length 13;
100.0%; Pred. No. 2.8e+02;
iive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PNA 3 inhibiting human and mammalian telomerase activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 13 BP; 1 A; 6 C; 2 G; 4 T; 0 U; 0 Other;
                          Piatyszek MA,
                                                                                                                                              Example 2; Col 37-38; 46pp; English.
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                                                                                                                    neurodegenerative diseases.
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(TEXA ) UNIV TEXAS SYSTEM.
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                                                   WPI; 2001-638024/73
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Best Local Similarity
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modified_base
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                          Shay JW,
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The present invention relates to peptide nucleic acids (PNAB), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in mammalian cells by hybridising to the RNA component of mammalian cells by hybridising to the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polynucleotide having the human telomerase RNA component (hTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the HTR gene RFLP pattern. The PNA can be further used for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmaceuticals (such as antinooplastic or cytostatic agents) can be used for treating nephasis, hymen immunodeficiency virus (HTV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomerame etabolism or telomerase activity. The present sequence represents one of the PNA sequences of the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases.
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                                                                                                                                                                                                                                                                                                                              Norton JC;
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                                                                                                                                                                                                                                                                                                                              Corey DR,
                                                                                                                                                                                                                                                                                                                              Piatyszek MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 2; Col 37-38; 46pp; English.
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                                                                                                                                             96US-00630019
97US-00838545
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                                                                                                                                                                                                                                                         (TEXA ) UNIV TEXAS SYSTEM
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Best Local Similarity 100.
Matches 13; Conservative
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                                                                                                                                                                                                                                                                                                                              Wright WE,
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                                                                     08-JUL-1999;
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09-APR-1997;
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25-SEP-2001
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The present invention relates to peptide nucleic acids (FNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in cammalian cells by hybridising to the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to denect and/or quantitate polynucleoride having the human telomerase RNA component (hTR) sequence, as well as in forensic identification of cindividuals, such as paternity testing or identification of individuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the hTR gene RFLP pattern. The PNA can be further used for treating or preventing cancer, inflammation, lymphoppicliferative diseases, autoinmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmaceuticals (such as antineophastic or cytostatic agents) can be used for treating neoplasia, hyperplasia, human immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomere metabolism or telomerase activity. The present sequence represents one of the PNA sequences of the invention. Note: The present sequence represents SED ID No 1 but is shown
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/*tag= a
//note= "This sequence is a peptide nucleic acid, i.e. it
contains a polyamide backbone instead of a deoxyribose
backbone"
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100.0%; Pred. No. 2.8e+02;
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97US-00838545.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmaceuticals (such as antineoplastic or cytostatic agents) can be used for treating neoplasia, hyperplasia, human immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomere metabolism or telomerase
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/label= OTHER
/note= "Phosphorothioate internucleotide linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                               New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or
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immunosuppressive; phosphorothioate; ss
                                                                                 Location/Qualifiers
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97US-00838545.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13; Conservative
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Matches 13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      present invention
                                                                                   Key
modified base
                                  sapiens
                                                                                                                                                                                                                                                  08-JUL-1999;
                                                                                                                                                                                                                                                                                   09-APR-1996;
                                                                                                                                                                                                                                                                                                   09-APR-1997;
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                                                                                                                                                                                    US6294650-B1
                                                                                                                                                                                                                   25-SEP-2001.
                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                    Shay JW,
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ID AAS154
XX AC AAS154
XX I4-FEE
XX DE PNA 2
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                                    Homo
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2'-arabino-fluorooligonucleotide; phosphoramidate; telomerase; inhibitor; infection; cancer; diagnosis; therapy; cytostatic; virucide; antisense;

/mod_base= OTHER /note= "2'-arabino-fluoronucleosides"

Location/Qualifiers

2. .13 /*tag= a

modified_base

Key

antigene; ss

Synthetic

/mod_base= OTHER /note= "phosphoramidate linkage"

/*tag= b

modified base

19-JAN-2001; 2001WO-US001918. 21-JAN-2000; 2000US-0178248P.

WO200153307-A1.

26-JUL-2001

Gryaznov S, Schultz RG; WPI; 2001-589652/66.

(GERO-) GERON CORP.

Phosphoramidate-linked 2'-arabino-fluorooligonucleotide.

(first entry)

26-NOV-2001

AAH26730;

AAH26730 standard; DNA; 13

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The present invention relates to peptide nucleic acids (PNAs), comprising a sequence of 6-25 nucleobases, that inhibit telomerase activity in mammalian cells by hybridising to the RNA component of mammalian telomerase. The PNAs are useful as probes to detect the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polynucleotide having the human telomerase RNA component (hTR) sequence, as well as in forensic identification of individuals, such as paternity testing or identification of criminal suspects or unknown descendants based on the HTR gene RFLP pattern. The PNA can be further used for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmaceuticals (such as attended plastic or cytostatic agents) can be used for treating neoplastic or cytostatic agents) can be used for treating neoplastic cor cytostatic agents) can be used for treating neoplastic cor cytostatic agents) and associated pathologies, and other diseases characterised by abnormal telomere metabolism or telomerase equivers.
                                                                                                                                                                                             /*tag= a
//note= "This sequence is a peptide nucleic acid, i.e. it
contains a polyamide backbone instead of a deoxyribose
backbone"
           Mammalian; peptide nucleic acid; probe; forensic; paternity testing; human telomeraes RNA component; hTR gene RFIP pattern; cancer; inflammation; lymphoproliferative disease; autoimmune disease; neurodegenerative disease; neoplasia; hyperplasia; HIV; AIDS; human immunodeficiency virus; acquired immunodeficiency virus; acquired immunodeficiency syndrome; telomere metabolism; mutant; cytostatic; atti-inflammatory; immunosuppressive; polyamide backbone; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or
                                                                                                                                                                                                                                                                                                                                                                                                                   Norton JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2.9%; Score 13; DB 1; Length 13;
100.0%; Pred. No. 2.8e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 13 BP; 8 A; 1 C; 2 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                   Corey DR,
                                                                                                                                                                                                                                                                                                                                                                                                                   Wright WE, Piatyszek MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 2; Col 37-38; 46pp; English.
                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                96US-00630019.
97US-00838545.
                                                                                                                                                                                                                                                                                                                      99US-00349532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                neurodegenerative diseases.
                                                                                                                                                                                                                                                                                                                                                                                        (TEXA ) UNIV TEXAS SYSTEM.
                                                                                                                                                                                            *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-638024/73
                                                                                                                                                                               modified base
                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                09-APR-1996;
09-APR-1997;
                                                                                                                                                                                                                                                               US6294650-B1
                                                                                                                                                                                                                                                                                                                      08-JUL-1999;
                                                                                                                                                                                                                                                                                          25-SEP-2001
                                                                                                                                        Synthetic
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                                                          The present sequence is that of a N3'-P5' 2'-arabino-fluoro phosphoramidate oligonucleotide that is complementary to telomerase RNA. The Oligonucleotide was used to assess the relative efficacy of novel 2'-arabino-fluoro phosphoramidate oligonucleotides and their 2'-ribo fluorooligonucleotide counterparts (see AAH26728-35) for the inhibition of telomerase activity. Novel phosphoramidate 2'-arabino-fluoroligonucleotides are generally more acid stable, more resistant to cellular proteases, and also show greater telomerase inhibition activity than 2'-ribose-fluoro phosphoramidates. They are therefore useful for treating cancer (claimed) and other diseases in which telomerase activity is present at abnormal levels, such as shearestoilferative or autoimmune diseases e.g. psoriasis, rheumacoid arthritis, immune system disordere requiring immunosuppression, and in the treatment of viral infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 13 BP; 5 A; 1 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.9%; Score 13; DB 1; Le
100.0%; Pred. No. 2.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
Example 6; Page 46; 61pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.08;
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Polynucleotides, used to detect and isolate nucleic acids, inhibit function of RNA and telomerase enzymes and to treat e.g. viral infections, contain 2'-arabino-fluoronucleoside(8) linked to

nucleoside(s).

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Gaps

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38 TTTTTTGTCTAAC 50 13; Conservative

Best Local Similarity Matches 13; Conserv

Query Match

TTTTTGTCTAAC 1

13

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RESULT 383 AAH26730/c

RESULT 384 AAH26734/c

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Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma; breast; prostate; colon; mixed cell leukaemia; Hodgkin's disease; fertility; inflammatory condition; tumour; cancer; veterinary; immunosuppression; telomerase inhibitor; ss.
                                                                        Human telomerase polynucleotide inhibitor #18
                                                                                                                                                                                                                                                                                                                               30-MAR-2001; 2001WO-US010476.
                                                                                                                                                                                                                                                                                                                                                       31-MAR-2000; 2000US-00540119
AAS15937 standard; DNA; 13
                                                                                                                                                                                                                                                                                                                                                                                                        Gryaznov SM, Pruzan R,
                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-656955/75
                                                                                                                                                                                                                                                                                                                                                                                (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                              WO200174136-A2
                                                                                                                                                                                                    Key
modified_base
                                                                                                                                                               sapiens.
                                                27-FEB-2002
                                                                                                                                                                                                                                                                                                      11-0CT-2001.
                                                                                                                                                                             Synthetic
oligonucleotide that is complementary to telomerase RNA. The arabino-leotide was used to assess the relative efficacy of novel 2'-arabino-luoro phosphoramidate oligonucleotides and their 2'-ribose fluorooligonucleotide counterparts (see AAH26728-35) for the inhibition of telomerase activity. Novel phosphoramidate 2'-arabino-luorooligonucleotides are generally more acid stable, more resistant to cellular proteases, and also show greater telomerase inhibition activity than 2'-ribose-fluoro phosphoramidates. They are therefore useful for treating cancer (claimed) and other diseases in which telomerase activity is present at abnormal levels, such as hyperproliferative or autoimmune diseases e.g. psoriaeis, rheumacoid arthritis, immune system disorders requiring immunosuppression, and in the treatment of viral infection
                                                                                                 2'-ribose-fluorooligonucleotide; phosphoramidate; telomerase; inhibitor;
                                                                                                              infection; cancer; diagnosis; therapy; cytostatic; virucide; antisense; antigene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Polynucleotides, used to detect and isolate nucleic acids, inhibit function of RNA and telomerase enzymes and to treat e.g. viral infections, contain 2'-arabino-fluoronucleoside(s) linked to nucleoside(s)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence is that of a 2'-ribose-fluoro phosphoramidate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.9%; Score 13; DB 1; Length 13; 00.0%; Pred. No. 2.8e+02;
                                                                        Phosphoramidate-linked 2'-ribose-fluorooligonucleotide.
                                                                                                                                                                                                                            /note= "2'-ribose-fluoronucleosides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 13 BP; 5 A; 1 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                              /note= "phosphoramidate linkage"
                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 6; Page 46; 61pp; English.
                                                                                                                                                                                                      /*tag= a
/mod_base= OTHER
                                                                                                                                                                                                                                                                  /mod_base= OTHER
ВР
                                                                                                                                                                                                                                                                                                                                                        19-JAN-2001; 2001WO-US001918.
                                                                                                                                                                                                                                                                                                                                                                                 21-JAN-2000; 2000US-0178248P.
AAH26734 standard; DNA; 13
                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Gryaznov S, Schultz RG;
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/*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                         (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                       WQ200153307-A1
                                                                                                                                                                                                                                           modified base
                                                                                                                                                                                        modified base
                                                  26-NOV-2001
                                                                                                                                                                                                                                                                                                                               26-JUL-2001.
                                                                                                                                                   Synthetic
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Weinrich SL;

/mod_base= OTHER /note= "N3'-P5' phosphoramidate linkages"

Location/Qualifiers

(first entry)

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1. .13 /*tag=

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                                                                                                                                                                                                                                                               The invention relates to polynucleotide inhibitors (I) and methods for inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity and proliferation of a telomerase positive cell, and in manufacturing a medicament for inhibiting telomerase activity in a manufacturing a medicament for inhibiting telomerase activity in a cell and in treating telomerase-mediated condition or disease, such as adenocarcinoma of breast, prostate or colon, mixed cell leukaemia, Hodgkin's disease, fertility and inflammatory conditions. (I) are also useful in treating a tumour or in manufacturing a medicament for the treatment of tumour. The polynucleotide inhibitors may also be used in diagnostic assays for detecting RNA or DNA. Inhibition of telomerase activity in cells in vivo is useful in prophylactic and therapeutic methods of treating cancer and other disorders involving inappropriate expression of telomerase, and in treating veterinary proliferative diseases. Inhibition of telomerase in haematopoietic stem cells is useful
                  or
New polynucleotide useful for inhibiting telomerase activity in cells, of for treating telomerase-mediated condition or disease, such as cancers, tumors, Hodgkin's disease, or inflammatory conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            branches of the immune system. The present sequence represents human telomerase polynucleotide inhibitor #18, as described in the method of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       represents human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             for immunosuppression and for selectively down-regulating specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ;
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1 Similarity 100.0%; Pred. No. 2.8e+02;
13; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 13 BP; 0 A; 1 C; 2 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                              Example 3; Page 32; 48pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            164 AGCAAACAAAAA 176
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Best Local Similarity
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RESULT 386

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Gaps

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0; Indels

100.0%; Pred. nv.

Best Local Similarity

Matches

Query Match

TIGICIAACCCIA 54 13; Conservative

42 13

TrGrcraaccera 1

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RESULT 385 AAS15937/c

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AAS15926,
                                                   The invention relates to polynucleotide inhibitors (I) and methods for inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity and proliferation of a telomerase positive cell, and in manufacturing a medicament for inhibiting telomerase activity in a cell and in treating telomerase-mediated condition or disease, such as and in treating telomerase-mediated condition or disease, such as and in treating telomerase mediated condition or disease, such as Hodgkin's disease, fertility and inflammatory conditions. (I) are also useful in treating a tumour or in manufacturing a medicament for the treatment of tumour. The polynucleotide inhibitors may also be used in treatment of tumour. The polynucleotide inhibitors and therapeutic methods of treating cancer and other disorders involving inappropriate contivity in cells in vivo is useful in prophylactic and therapeutic methods of treating cancer and other disorders involving inappropriate diseases. Inhibition of telomerase in haematopoletic stem cells is useful for thmunosuppression and for selectively down-regulating specific.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   branches of the immune system. The present sequence represents human telomerase polynucleotide inhibitor \#2, as described in the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New polynucleotide useful for inhibiting telomerase activity in cells, for treating telomerase-mediated condition or disease, such as cancers, tumors, Hodgkin's disease, or inflammatory conditions.
                                                                                                           Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma; breast; prostate; colon; mixed cell leukaemia; Hodgkin's disease; fertility; inflammatory condition; tumour; cancer; veterinary; immunosuppression; telomerase inhibitor; ss.
                                                                                                                                                                                                                            1. .13
/*tag= a
/mod_base= OTHER
/note= "N3'-P5' phosphoramidate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 13 BP; 4 A; 2 C; 6 G; 1 T; 0 U; 0 Other;
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100.0%; Pred. No. 2.8e+02;
ive 0; Mismatches 0;
                                                                                      Human telomerase polynucleotide inhibitor #2.
                                                                                                                                                                                                                                                                                                                                                                                                                    Weinrich SL;
                                                                                                                                                                                                                 Location/Qualifiers
Claim 8; Page 36; 48pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
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Matches
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The invention relates to polynucleotide inhibitors (I) and methods for inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity and proliferation of a telomerase positive cell, and in manufacturing a medicament for inhibiting telomerase activity in a cell and in treating telomerase. Fertility and inflammatory conditions. (I) are also the definity a timour or in manufacturing a medicament for the treatment of tumour. The polynucleotide inhibitors may also be used in treatment of tumour or in manufacturing a medicament for the treatment of tumour. The polynucleotide inhibitors may also be used in diagnostic assays for detecting RNA or DNA. Inhibition of telomerase activity in cells in vivo is useful in prophylactic and therapeutic methods of treating cancer and other disorders involving inappropriate expression of telomerase, and in treating veterinary proliferative concludation of telomerase in the mematopoietic stem cells is useful to branches of the immune system. The present sequence represents human telomerase polynucleotide inhibitor #7, as described in the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ô
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New polynucleotide useful for inhibiting telomerase activity in cells, or for treating telomerase-mediated condition or disease, such as cancers, tumors, Hodgkin's disease, or inflammatory conditions.
                                                                                                                                                                                                                                                                  Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma;
breast; prostate; oclon; mixed cell leukaemia; Hodgkin; a disease;
fertility; inflammatory condition; tumour; cancer; veterinary;
immunosuppression; telomerase inhibitor; ss.
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Pred. No. 2.8e+02;
                                                                                                                                                                                                         Human telomerase polynucleotide inhibitor #7.
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100.0%; Pred
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                         ВР
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926/c
AAS15926 standard; DNA; 13
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                                                                                                                                               (first entry)
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Local Sim
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                                                                               AAS15926;
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143 GCCTTCCACCGTT 155 Conservative

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13;

13 GCCTTCCACCGTT 1

Human telomerase polynucleotide inhibitor #16

(first entry)

27-FEB-2002

AAS15935;

BP.

AAS15935 standard; DNA; 13

AAS15935/

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inhibiting telomerase activity. (1) are useful in inhibiting telomerase activity and proliferation of a telomerase positive cell, and in manufacturing a medicament for inhibiting telomerase activity in a cell and in treating telomerase activity in a cell and in treating telomerase—mediated condition or disease, such as adenocarcinoma of breast, prostate or colon, mixed cell leukaemia, Hodgkin's disease, fertility and inflammatory conditions. (1) are also useful in treating a tumour or in manufacturing a medicament for the treatment of tumour. The polynucleotide inhibitors may also be used in diagnostic assays for detecting RNA or DNA. Inhibition of telomerase activity in cells in vivo is useful in prophylactic and therapeutic methods of treating cancer and other disorders involving inappropriate expression of telomerase, and in treating veterinary proliferative diseases. Inhibition of telomerase in hemanopoietic stem cells is useful for immunosuppression and for selectively down-regulating specific for immunosupression and for selectively down-regulating specific branches of the immune system. The present sequence represents human
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                                                                                                                                                                                                                                                                                                                                                                                     Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma; breast; prostate; colon; mixed cell leukaemia; Hodgkin's disease; fertility; inflammatory condition; tumour; cancer; veterinary; immunosuppression; telomerase inhibitor; ss.
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                                                                                                                                                                                                                                                                                                              Human telomerase polynucleotide inhibitor #11
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                                                                       AAS15930 standard; DNA; 13 BP
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/*tag= a
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modified_base
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RESULT 388
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The invention relates to polynucleotide inhibitors (I) and methods for inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity in a calmanifacturing a medicament for inhibiting telomerase activity in a cell and in treating telomerase-mediated condition or disease, such as adenocarcinoma of breast, prostate or colon, mixed cell leukaemia. CC Hodgkin's disease, fertility and inflammatory conditions. (I) are also useful in treating a tumour or in manufacturing a medicament for the treatment of tumour. The polynucleotide inhibitions may also be used in diagnostic assays for detecting RNA or DNA. Inhibition of telomerase activity in cells in vivo is useful in prophylactic and therapeutic methods of treating cancer and other disorders involving inappropriate expression of telomerase, and in treating veterinary proliferative consists and consists a
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                                                                                                                                                                                    Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma; breast; prostate; colon; mixed cell leukaemia; Hodgkin's disease; fertility; inflammatory condition; tumour; cancer; veterinary; immunosuppression; telomerase inhibitor; ss.
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/note= "N3'-P5' phosphoramidate linkages"
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/*tag= a
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"N3'-P5' F
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modified base
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Gaps ·.

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2.9%; Score 13; DB 1; Length 13; 00.0%; Pred. No. 2.8e+02;

154 TTCATTCTAGAGC 166

à a

TTCATTCTAGAGC 1

13

13; Conservative

Matches

Local Similarity

Query Match

13 AAACAAAAATGT

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The invention relates to polynucleotide inhibitors (I) and methods for inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity and proliferation of a telomerase positive cell, and in manufacturing a medicament for inhibiting telomerase activity in a call and in treating telomerase-mediated condition or disease. Such as adenocarcinoma of breast, prostate or colon, mixed cell leukaemia, Hodgkin's disease, fertility and inflammatory conditions. (I) are also useful in treating a tumour or in manufacturing a medicament for the treatment of tumour. The polynucleotide inhibitors may also be used in diagnostic assays for detecting RNA or DNA. Inhibition of telomerase activity in cells in vivo is useful in prophylactic and therapeutic methods of treating cancer and other disorders involving inappropriate expression of telomerase, and in treating veterinary proliferative diseases. Inhibition of telomerase in haematopoietic stem cells is useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  for immunosuppression and for selectively down-regulating specific branches of the immune system. The present sequence represents human telomerase polynucleotide inhibitor #3, as described in the method of the
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                                                                                                                                                                                  Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma;
breast; prostate; colon; mixed cell leukaemia; Hodgkin's disease;
fertility; inflammatory condition; tumour; cancer; veterinary;
immunosuppression; telomerase inhibitor; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       polynucleotide useful for inhibiting telomerase activity in cells,
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                                                                                                                                                  Human telomerase polynucleotide inhibitor #3
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AAS15922 Btandard; DNA; 13 BP.
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                                                                        AAS15922;
RESULT 390
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                 AAS15922
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The invention relates to polynucleotide inhibitors (I) and methods for inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity and proliferation of a telomerase positive cell, and in manufacturing a medicament for inhibiting telomerase activity in a cell and in treating telomerase. mediated condition or disease, such as adenocarcinoma of breast, prostate or colon, mixed cell leukaemia, Hodgkini's disease, fertility and inflammatory conditions. (I) are also useful in treating a tumour or in manufacturing a medicament for the treatment of tumour. The polynucleotide inhibitors may also be used in diagnostic assays for detecting RNA or DNA. Inhibition of telomerase activity in cells in vivo is useful in prophylactic and therapeutic methods of treating cancer and other disorders involving inappropriate expression of telomerase, and in treating veterinary proliferative confinements. Inhibition of telomerase in haematopoidtic stem cells is useful for immunosuppression and for selectively down-regulating specific confinements.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New polynucleotide useful for inhibiting telomerase activity in cells, or for treating telomerase-mediated condition or disease, such as cancers, tumors, Hodgkin's disease, or inflammatory conditions.
                                                                                                                                                                                                                                  Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma; breast; prostate; colon; mixed cell leukaemia; Hodgkin's disease; fertility; inflammatory condition; tumour; cancer; veterinary; immunosuppression; telomerase inhibitor; ss.
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/note= "N3'-P5' phosphoramidate linkages"
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                                                                                                                                                                                                  Human telomerase polynucleotide inhibitor #4.
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                                                                                       AAS15923 standard; DNA; 13
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/*tag= a
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Matches 13; Conserv
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                                                     RESULT 391
AAS15923/c
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2.9%; Score 13; DB 1; Length 13; 100.0%; Pred. No. 2.8e+02; tive 0; Mismatches 0; Indels

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The invention relates to polymucleotide inhibitors (I) and methods for inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity in a can activity and inhibiting telomerase activity in a cell and in reading telomerase—mediated condition or disease, such as adenocarcinoma of breast, prostate or colon, mixed cell leukaemia, chodykin's disease, fertility and inflammatory conditions. (I) are also useful in treating a tumour or in manufacturing a medicament for the treatment of tumour. The polymucleotide inhibitors may also be used in diagnostic assays for detecting RNA or DNA. Inhibition of telomerase activity in cells in vivo is useful in prophylactic and therapeutic methods of treating cancer and other disorders involving inappropriate cypression of telomerase, and in treating veterinary proliferative diseases. Inhibition of telomerase in haematopoietic stem cells is useful continuous continuous
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma; breast; prostate; colon; mixed cell leukaemia; Hodgkin's disease; fertility; inflammatory condition; tumour; cancer; veterinary; immunosuppression; telomerase inhibitor; ss.
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/note= "N3'-P5' phosphoramidate linkages"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human telomerase polynucleotide inhibitor #6
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                                                                                                                                                                                                                                                                                                                                                                                  BP
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                                                                                                                                                                                                                                                                                                                                                                                        AAS15925 standard; DNA; 13
CCGCCTTCCACCG 153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                         CCGCCTTCCACCG 1
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/*tag=
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modified_base
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The invention relates to polynuclectide inhibitors (I) and methods for inhibiting telomerase activity. (I) are useful in inhibiting telomerase activity and proliferation of a telomerase positive cell, and in cativity and proliferation of a telomerase positive cell, and in manufacturing a medicament for inhibiting telomerase activity in a cell and in treating telomerase. mediated condition or disease, such as adenocarcinoma of breast, prostate or colon, mixed cell leukaemia, Hodgkin's disease, fertility and inflammatory conditions. (I) are also useful in treating a tumour or in manufacturing a medicament for the treatment of tumour. The polynuclectide inhibitors may also be used in diagnostic assays for detecting RNA or DNA. Inhibition of telomerase activity in cells in vivo is useful in prophylactic and therapeutic methods of treating cancer and other disorders involving inappropriate expression of telomerase, and in treating veterinary proliferative diseases. Inhibition of telomerase in haematopoietic stem cells is useful for immunosuppression and for selectively down-regulating specific.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New polynucleotide useful for inhibiting telomerase activity in cells, or for treating telomerase-mediated condition or disease, such as cancers, tumors, Hodgkin's disease, or inflammatory conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   branches of the immune system. The present sequence represents human telomerase polynucleotide inhibitor #5, as described in the method of the
                                                                                                                                                                                                                                                                                        Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma; breast; prostate; colon; mixed cell leukaemia; Hodgkin's disease; fertility; inflammatory condition; tumour; cancer; veterinary; immunosuppression; telomerase inhibitor; ss.
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/mod_base= OTHER
/note= "N3'-P5' phosphoramidate linkages"
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                                                                                                                                                                                                                                                     Human telomerase polynucleotide inhibitor #5.
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                                                                                                                                    AAS15924 standard; DNA; 13
138 CTGCCGCCTTCCA 150
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                                                                                                RESULT 393
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2.9%; Score 13; DB 1; Length 13; 100.0%; Pred. No. 2.8e+02;

Query Match Best Local Similarity

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Gaps . 0

Query Match 2.9%; Score 13; DB 1; Length 13; Best Local Similarity 100.0%; Pred. No. 2.8e+02; Matches 13; Conservative 0; Mismatches 0; Indels

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The invention relates to polynucleotide inhibitors (1) and methods for inhibiting telomerase activity. (1) are useful in inhibiting telomerase activity. (1) are useful in inhibiting telomerase activity and proliferation of a telomerase positive cell, and in manufacturing a medicament for inhibiting telomerase activity in a cell and in treating telomerase-mediated condition or disease, such as adenorating telomerase-mediated condition or disease, such as adenorated fertility and inflammatory conditions. (1) are also useful in treating a tumour or in manufacturing a medicament for the treating a tumour or in manufacturing a medicament for the disposetic assays for detecting RNA or DNA. Inhibition of telomerase activity in cells in vivo is useful in prophylactic and therapeutic methods of treating cancer and other disorders involving inappropriate expression of telomerase, and in treating veterinary proliferative diseases. Inhibition of telomerase in haematopoietic stem cells is useful for selectively down-regularing specific branches of the immune system. The present sequence represents human telomerase polynucleotide inhibitor #19, as described in the method of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New polynuclectide useful for inhibiting telomerase activity in cells, for treating telomerase-mediated condition or disease, such as cancers, tumors, Hodgkin's disease, or inflammatory conditions.
                                                                                                                                                                                                                                                                                      Human; telomerase; hTR; cytostatic; anti-inflammatory; adenocarcinoma; breast; prostate; colon; mixed cell leukaemia; Hodgkin's disease; fertility; inflammatory condition; tumour; cancer; veterinary; immunosuppression; telomerase inhibitor; ss.
Gaps
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Indels
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                                                                                                                                                                                                                                                        Human telomerase polynucleotide inhibitor #19.
0; Mismatches
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                                139 TGCCGCCTTCCAC 151
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13; Conservative
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Matches
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AAS15938/0
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2.9%; Score 13; DB 1; Length 13;

Query Match

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and anti-gene diagnostic or therapeutic applications and may be used for treating telomerase-mediated conditions or diseases, such as hyperproliferative and autoimmune disorders, and for contraceptive
                                                                                                                                                                                                                                                                                                          Thiophosphoramidate oligonucleotide; virucide; cytostatic; immunosuppressive; contraceptive; RNA inhibitor; telomerase inhibitor; antisense therapy; viral infection; cancer; hyperproliferative disorder; autoimmune disorder; se.
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                             Gaps
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.00.0%; Pred. No. 2.8e+02;
.ve 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                             Thiophosphoramidate oligonucleotide, SEQ ID NO: 2.
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          Pred. No. 2.8e+02; ; Mismatches 0;
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100.08; PL
                                                                                                                                                                              BP
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99US-0160444P.
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                                                             161 TAGAGCAAACAAA 173
                                                                                                                                                                                                                                              (first entry)
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                                Conservative
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             Local Similarity
les 13; Conserv
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tes 13; Conserv
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             Best Loca
Matches
                                                                                                                                              RESULT 395
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Oligonucleotide #1 used in conjugates of the invention.
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93WO-US009297.
95US-00403888.
99US-00299058.
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                                                                                                                                           23-MAR-2001; 2001US-0278322P.
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es 13; Conservative
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                                                                                                                                                                                                                                                        telomerase inhibitors.
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                                                                        WO200277184-A2
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29-SEP-1993;
12-JUN-1995;
23-APR-1999;
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Matches
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                                                                                                                    Thiophosphoramidate oligonucleotide, virucide, cytostatic, immunosuppressive, contraceptive, RNA inhibitor, telomerase inhibitor; antisense therapy, viral infection, cancer, hyperproliferative disorder;
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                                                                                                Thiophosphoramidate oligonucleotide, SEQ ID NO: 10.
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                          AAF81195 standard; DNA; 13 BP
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99US-0160444P.
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                                                                                                                                                                                                                                                                                                                                              Gryaznov S, Pongracz K,
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                                                                                                                                                            autoimmune disorder; ss
                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-265967/27
                                                                                                                                                                                                                                                                                                                     (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                           WO200118015-A1
                                                                                                                                                                                                                                                                                 10-SEP-1999;
19-OCT-1999;
                                                                         30-MAY-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13;
                                                                                                                                                                                                                                   15-MAR-2001
                                                                                                                                                                                   Synthetic.
                                                 AAF81195;
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  RESULT 396
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             AADS0105/c
ID AADS0
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AC AADS0
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DT 24-M
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New oligonucleotide conjugate useful in the treatment of e.g. cancers are
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Match phosphorothioate modified 2'-0-methyl RNA G-core oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chiang M, Brown-Driver VL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .
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Medicine; telomerase; tumour; cancer; leukaemia; lymphoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 13; DB 1; Length 13;
Pred. No. 2.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 13 BP; 5 A; 1 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Morin GB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Anderson KP, Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tolman RL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 12; Page 37; 45pp; English.
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99US-00299058
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23-APR-1999;
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                                                                                               Ecker DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     46
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABZ58497;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Reducing
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                                                                                                                                                                                         The invention relates to a novel chemically modified oligonucleotide having no more than about 27 nucleic acid base units. The oligonucleotide modulates mammalian telomere length. The chemically modified oligonucleotide of the invention may be useful for modulating the telomere length of a mammalian chromosome, inhibiting the division of a mammalian cell or modulating the effects of aging of a mammalian cell. The oligonucleotides may also useful for treating diseases associated with abnormal telomere length such as aging and hyperproliferative conditions including cancer. The current sequence is that of the "match" phosphorothhoate modified 2'-O-methyl RNA G-core oligonucleotide of the invention.

    1. 13
    *tag= b
    *mod_base= OTHER
    *note= "OTHER = Optionally phosphorothioate backbone and 2'-O-methyl sugar modification"

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/mod_base= mSc, OTHER
/mod_bas
                                                                           New chemically modified oligonucleotides, useful for modulating telomere length of a mammalian chromosome, inhibiting the division of a malignant mammalian cell, or modulating the effects of aging of a mammalian cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        telomere length; aging; hyperproliferative condition; cancer; ss; PNA; peptide nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               note = "OTHER = Optionally linked to Gly residue
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.9%; Score 13; DB 1; Length 13; 100.0%; Pred. No. 2.8e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Match 2'-O-methyl oligonucleotide / peptide nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 13 BP; 3 A; 1 C; 5 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                        Example 5; Page 6; 10pp; English.
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/mod base= OTHER
    Wyatt JR
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ADB68046 standard; DNA; 13 BP.
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93WO-US009297.
95US-00403888.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 100.
Marches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       46 CTAACCCTAACTG 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CTAACCCTAACTG 1
  Vickers TA,
                                        WPI; 2003-606442/57.
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12-JUN-1995;
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    Ecker DJ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mammalian hair growth comprises applying telomerase inhibitor to
                                                                                                                                                                                                   New chemically modified oligonucleotides, useful for modulating telomere length of a mammalian chromosome, inhibiting the division of a malignant mammalian cell, or modulating the effects of aging of a mammalian cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence is that of telomerase inhibitor VI, an example a telomerase inhibitor useful in the method of the invention for reducing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gарв
                                                         Brown-Driver VL;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
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                                                             ΣÌ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 13 BP; 3 A; 1 C; 5 G; 4 T; 0 U; 0 Other;
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Pred. No. 2.8e+02;
                                                             Chiang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bennett CF,
                                                   A. Wyatt JR;
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                                                                                                                                                                                                                                                                                                                          Example 5; Page 6; 10pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        497/c
ABZ58497 standard; RNA; 13
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                                                      RC, Anderson K
I, Vickers TA,
(ISIS-) ISIS PHARM INC
                                                                                                                                            WPI; 2003-606442/57.
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Best Local Similarity
Matches 13; Conserv
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The invention relates to a novel compound having telomerase inhibition activity, comprising a nucleobase (A) and oligonucleotide (O) containing a sequence of 2-11 nucleotides exactly complementary to a sequence within the template region of human telomerase RNA, linked through a linker (L). A compound of the invention has cytostatic activity. The compound is useful for inhibiting the activity of telomerase enzyme, which involves contacting the telomerase with the compound. The compound of the invention is also useful for inhibiting the proliferation of the cinvention is also useful for inhibiting the proliferation of the cinvention cell: The compound is useful as a medicine for retaing cancer. The compound is useful for inhibiting or reducing treating cancer cell: The compound is useful as a medicine for the commerase activity and/or proliferation of cells having telomerase activity e.g., tumour cells. The telomerase-positive tumour cells are from cancer of the skin, connective tissue, adipose, breast, lung, stomach, pancreas, ovary, cervix, uterus, kidney, bladder, colon, prostate, central nervous system (CNS), retina and circulating tumours
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unwanted mammalian (human) hair growth. A composition comprising the telomerase inhibitor is applied to the skin to reduce hair growth. The telomerase inhibitor acts by reducing telomerase levels in hair follicles, reducing telomerase mRNA expression or by promoting the erosion of telomeric DNA. It can be used to reduce hair growth of a woman with hirsutism or to reduce androgen stimulated hair growth (both claimed). A composition comprising a telomerase inhibitor reduces hair growth by at least 15% (preferably 20%) when tested in the Golden Syrian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel covalent oligonucleotide conjugates comprising nucleobase and oligonucleotide exactly complementary to sequence within template region of human telomerase RNA, linked through linker, useful as medicine for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ss; telomerase; hTR; template region; cytostatic; telomerase inhibition;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Telomerase template region complementary oligonucleotide SEQ ID NO:3.
                                                                                                                                                                                                                                                                                      Gaps
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Pred. No. 2.8e+02;
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                                                                                                                                                                                                       Sequence 13 BP; 3 A; 1 C; 5 G; 0 T; 4 U; 0 Other;
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                                                                                                                                                                                                                                2.9%; Scor.
100.0%; Pred. No. 2...
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP.
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                                                                                                                                                                                                                                                                                      13; Conservative
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                                                                                                                                                                                                                                              Query Match
Best Local Similarity
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                                                                                                                                                                    Hamster assay
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-JUL-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 401
                                                                                                                                                                                                                                                                                      Matches
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/note= "OTHER = Optionally attached to C3H711CONH group"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a novel method for manufacturing a labelled nucleic acid which comprises introducing an 11C to the transduction region and contacting the nucleic acid fragment and 11C compound, where the nucleic acid fragment is a polynucleic acid or peptide nucleic acid. The method of the invention may be useful for manufacturing a labelled nucleic acid and provides sensitive detection of a target mRNA, such neuroglia myofibril acidic protein (GFAB). The current sequence is that of the labelled nucleic acid/l1C-related PNA-antisense oligo 2 of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Manufacturing labeled nucleic acid, by introducing carbon-11 to transduction region, and contacting nucleic acid fragment and carbon-11
                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
(such as leukaemia and lymphoma). The present sequence represents an oligonucleotide of the invention.
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                                                                                                                                                                                                                                                                                                                                                                              labelled nucleic acid manufacture; 11C; sensitive detection;
                                                                           2.9%; Score 13; DB 1; Length 13; 100.0%; Pred. No. 2.8e+02; iive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.9%; Score 13; DB 1; Length 13; 100.0%; Pred. No. 2.8e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                Labelled nucleic acid/11C-related PNA-antisense oligo 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 13 BP; 3 A; 1 C; 5 G; 4 T; 0 U; 0 Other;
                                                Seguence 13 BP; 5 A; 1 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 2; Fig 7; 17pp; Japanese.
                                                                                                                                                                                                                                                     BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (NIKL ) JAPAN STEEL WORKS LTD.
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                                                                                                                                                                                                                                                     ADO21607 standard; DNA; 13
                                                                                                                                                                                                                                                                                                                  (first entry)
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                                                                                                                                            42 TTGTCTAACCCTA 54
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es 13; Conservative
                                                                                                             13; Conservative
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                                                                                Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                          Unidentified
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16-MAR-2004; 2004WO-US008129. 17-MAR-2003; 2003US-0455705P. (MEDI-) MEDICAL COLLEGE OHIO Example; SEQ ID NO 3; 99pp; English.

WPI; 2004-690896/67.

Ratnam M;

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The invention relates to a novel method for diagnosing and treating a patient (suspected of) having a solid tumour that produces folate receptor athoral (Re-alpha), comprising administering a folate receptor receptor athoral (Re-alpha), comprising administering a folate receptor applies (Re-alpha) inducer comprising at mistoring a folate receptor applies inducer increases levels of the receptor on plasma membranes of tumour cells and in at least one type of body fluid. Also claimed as part of the invention are: a method for diagnosing or therapeutically treating a patient comprising activating the folate receptor alpha (RR-alpha) gene in the patient using at lest one of an agonist or antagonist of cestrogen receptor (ER), progesterone receptor (FR), androgen receptor (RR) or glucocorticoid receptor (RR); a method for substantially increasing folate receptor agent in a manner or tumour using at least one steroid receptor agent in a manner selective to FR-alpha-postive tissues; and finally a method to enhance sensitivity of whole body imaging and the therapeutic efficacy of FR-alpha; and a serum marker for diagnostic exceptor agent that binds to cancers during and after treatment comprising and monitoring of cancers during and after treatment comprising lucescent conjugates of folate or antibodies specific for the folate receptor. Method of the invention has cytostatic activity. The method of having, at least one solid tumour. The present sequence represents a human G/C-rich (Spl binding)
                                                                                                                                                                                              Diagnosing and treating a patient with a solid tumor that produces folate receptor alpha, comprises administering a folate receptor alpha inducer comprising a steroid receptor agent, optionally with a histone deacetylase inhibitor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 13 BP; 1 A; 0 C; 10 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a novel method for manufacturing a labelled nucleic acid which comprises introducing an 11C to the transduction region and contacting the nucleic acid fragment and 11C compound, where the nucleic acid fragment is a polynucleic acid or peptide nucleic acid. The method of the invention may be useful for manufacturing a labelled nucleic acid and provides sensitive detection of a target mRNA, such an energalia myofibril acidic protein (GFAP). The current sequence is that of the labelled nucleic acid/11C-related PNA-antisense oligo 1 of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Manufacturing labeled nucleic acid, by introducing carbon-11 to transduction region, and contacting nucleic acid fragment and carbon-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                            abelled nucleic acid manufacture; 11C; sensitive detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.9%; Score 13; DB 1; Length 13; 100.0%; Pred. No. 2.8e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FR-alpha; G/C rich; solid tumour; folate receptor alpha; steroid receptor; cancer; cytostatic; ds; Spl binding.
                                                                                                                                                          Labelled nucleic acid/11C-related PNA-antisense oligo 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 13 BP; 5 A; 1 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human G/C-rich (Sp1 binding) element SEQ ID NO:3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 2; Fig 7; 17pp; Japanese.
                                                 ADO21606 standard; DNA; 13 BP
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                                                                                                                        (first entry)
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13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13 TTGTCTAACCCTA 1
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                                                                                                                                                                                                                 PNA-antisense; ss
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                                                                                                                                                                                                                                                                                       JP2004123542-A.
                                                                                                                                                                                                                                                   Unidentified
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                                                                                                                     15-JUL-2004
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Best Local S:
Matches 13,
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                RESULT 403
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Human, c-raf, A-raf, B-raf, hammerhead ribozyme, hairpin ribozyme,
target, substrate; catalyst, modulation, expression, Raf gene, delivery,
screening, identification, synthesis, deprotection, purification, cancer,
inflammation, psoriasis, non-hepatic ascites, infection, genetic drift;
restenosis, rheumatoid arthritis, ss.
                                                                                                                                                                             Human B-raf target sequence nucleotide position 2205.
                                                                              BP
                                                                              AAV93815 standard; RNA; 14
                                                                                                                                             18-FEB-1999 (first entry)
GGGAGGGGTGGTG 13
                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                              AAV93815;
                                                RESULT 405
                                                            AAV93815
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Gарв ö

2.9%; Score 13; DB 1; Length 13; 100.0%; Pred. No. 2.8e+02; ive 0; Mismatches 0; Indels

21 GGGAGGGGTGGTG 33

Local Similarity 100. 1es 13, Conservative

element

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The present invention relates to novel 2',5'-oligoadenylic acid analogs and their pharmacologically- acceptable salts. The analogs are stable with superior antitumour and antiviral activity and so are useful in cancer or antiviral therapy e.g. as antisense drugs. The present sequence was used to illustrate the invention.
                                                                                                                                         /mod_base= OTHER
/note= "2',4'-oxyethylene linkage in the sugar residues
and phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Stable 2',5'-oligoadenylic acid analogs containing natural and modified nucleic acid units as well as unusual phosphate groups with excellent activity particularly antitumor, applicable in cancer or antiviral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Galanin receptor; GALR1; human; single nucleotide polymorphism; SNP; drug discovery; haplotyping; infectious diarrhoea; growth hormone deficiency; allele-specific oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Galanin receptor gene GALR1 allele-specific oligonucleotide #46
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                                                                                                                                                                                                                                                                /mod_base= OTHER
/note= "G-hydroxyethyl phosphate"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 14 BP; 5 A; 1 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 111; 220pp; Japanese.
                                                                    Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                     19-NOV-2003; 2003WO-JP014748.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-NOV-2002; 2002JP-00334731
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AAS98334 standard; DNA; 15
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Best Local Similarity 100.
Matches 13; Conservative
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                                                                                                                    '*tag≃
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (SANY ) SANKYO CO LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Koizumi M, Morita K;
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                                                                                           modified base
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                      Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Action has been developed for the identification of a nucleic acid capable of modulating a process in a biological system. The method capable of modulating a process in a biological system. The method capable of introducing into the system a random library of nucleic acid catalysts (NAC) having a substrate binding domain (SBD), comprising a random sequence, and a catalytic domain (CD); and (b) identifying NBC in systems where modulation has occurred and/or determining the sequence of at least part of the SBDs in such systems. Nucleic acid molecules with cendonuclease activity and catalytic activity, from the present invention. The are used to modulate gene expression in plant and mammalian cells and to cleave target nucleic acid, particularly for treating systemic diseases caused by specific RNA, e.g. cancer, inflammation, psoriasis, non-hepbatic ascites and infection. They may also be used to detect genetic drift and mutations in diseased cells and to determine c-raf RNA. Specifically NACs with RNA-cleaving activity that modulate expression of the Raf gene, are used to treat cancer, restenosis, psoriasis or rheumatoid arthritis, or seed to treat cancer, restenosis, psoriasis or rheumatoid arthritis, or of sugar/phosphate modifications increases stability against nuclease and activity. AAV99922 to AAV93877 represent NACs that can be used in the method, specifically for modulating the expression of a Raf gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Identifying new catalytic nucleic acid that modulates selected processes - especially ribozymes that cleave Raf RNA for treating cancer, restenosis, and also new ribozymes and modified nucleoside triphosphates used as antiviral agents and synthons.
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                                                                                                                                                                                                                                                                                                                                                                                                              Bellon L;
Burgin A;
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                                                                                                                                                                                                                                                                                                                                                                                                         nic J, Reynolds M, Kisich K,
Mcswiggen JA, Karpeisky A,
T, Beaudry A, Sweedler D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 14 BP; 1 A; 5 C; 6 G; 0 T; 2 U; 0 Other;
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                                                                                                                                                                                      970S-0051718P
970S-0056808P.
970S-0061321P.
970S-0061324P.
970S-0064866P.
                                                                                                                                           97US-0046059P.
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                                                                                                                                                                                                                                                                                                                   97US-0068212P
                                                                                                98WO-US009249
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                                                                                                                                                                                                                                                                                                                                                                   (RIBO-) RIBOZYME PHARM INC.
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Parry T, Beigelman L, M
Thompson J, Workman CT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AGCUGCUGGCCCG 14
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  WO9850530-A2
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                                                                                                05-MAY-1998;
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22-AUG-1997
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05-NOV-1997
                                                12-NOV-1998
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  NAMES OF COLOR COL
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Gaps

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(GENA-) GENAISSANCE PHARM INC

03-JAN-2001; 2001WO-US000092. 03-JAN-2001; 2001WO-US000092

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The invention relates to genotyping human galanin receptor (GALRI) gene of an individual, involving determining for the two copies of the GALRI of an individual, in the indentity of the nucleotide pair at come or more polymorphic sites. The method is useful for determining whether an individual has a haplotype or haplotype pairs defined in the specification. This is useful for improving the efficacy and reliability of several steps in the discovery and development of drugs for treating diseases associated with GALRI activity, e.g., infectious diarrhoea and growth hormone deficiency, to validate GALRI as a candidate agent for treating a specific condition or disease predicted to be associated with GALRI activity, and in the design of clinical trials of candidate drugs for treating a specific condition or disease predicted to be associated with GALRI activity. The method is useful to compounds the expression and function of GALRI, and in sypressing GALRI protein for cargoring GALRI activity. The method is useful for studying expression and function of GALRI, and in sypressing GALRI protein for use in screening for candidate drugs to treat diseases related to GALRI activity. The polymucleotide or variant is useful for studying expression or the biological activity of GALRI as well as on the binding the affinity of candidate drugs teresonening and testing of drugs crayistion on the biological activity of GALRI as well as on the binding caffinity of candidate drugs teresonening and testing of the cypersent human GALRI gene and growth hormone insufficiency. ASS98289- ASS98408 represent human GALRI gene allele-specific oligonucleotides used to invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                           Genotyping human galanin receptor gene of an individual for determining haplotype of an individual, involves determining the identity of nucleotide pair at specific polymorphic sites for two copies of the gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lecithin-cholesterol acyltransferase; LCAT; Norum disease; gene therapy; fish-eye disease; atherosclerotic cardiovascular disease; forensic; population diversity; anthropological lineage; paternity testing; human; polymorphism; allake-specific oligonucleotide; ASO; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                         Denton RR, Nandabalan K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 13; DB 1; Length 15;
Pred. No. 3.4e+02;
1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human LCAT gene polymorphism detection ASO primer #16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 15 BP; 2 A; 3 C; 9 G; 0 T; 0 U; 1 Other;
                                                                                                         Choi JY,
                                                                                                                                                                                                                                                                               Claim 16; Page 15; 99pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABK97507 standard; DNA; 15 BP.
                                                               (GENA-) GENAISSANCE PHARM INC
                    14-APR-2000; 2000US-0197838P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 2.9%;
Best Local Similarity 86.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              82 TTGCTCCCGCGCGC 96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15 TYCCTCCCGCGCGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13; Conservative
                                                                                                         Chew A,
                                                                                                                                                    WPI; 2002-066341/09.
                                                                                                         Bentivegna SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200253575-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABK97507;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 408
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The present invention relates to a new polynucleotide comprising a nucleotide sequence which is a polymorphic variant of a reference sequence which is a polymorphic variant of a reference sequence for lecithin-cholesterol acyltransferase (LCAT). The invention is useful for identifying an association between a trait (preferably a clinical response to drug targeting LCAT) and at least one genotype or haplotype of LCAT gene. The method of the invention has applicability in developing diagnostic tests and therapeutic treatments for Norum disease, is lab-eye disease and atherosclerotic cardiovascular disease. The haplotyping and genotyping methods are useful for studying population diversity, anthropological lineage, the significance of diversity and for identifying association between the LCAT sentence applications and for identifying section between the LCAT maplotypes greent in individuals are useful in the development of drugs targeting LCAT. For example, determining the frequency of individual LCAT haplotypes in a individual are useful in the development of drugs targeting the LCAT individual ECAT. For example, determining the frequency of individual LCAT haplotypes in a completion with a specific disease, e.g. Norum disease, will facilitate the development of drugs targeting the LCAT isoform(8) that are most frequent in that disease population. The present nucleic acid sequence of example and sequence of sequence of solection (ABKSY192-ABKSPIS) of allele specific contents on the contents on the present nucleic acid sequence of a collection (ABKSY192-ABKSPIS) of allele specific
                                                                                                                                                                                                                                    Novel isolated polymorphic variant polynucleotide of lecithin-cholesterol acyltransferase gene, useful for studying expression and biological function of the gene, and for therapeutic, diagnostic or forensic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               that were used in the invention to detect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 human; cholesteryl ester transfer protein; CETP;
single nucleotide polymorphism; SNP; drug screening; atherosclerosis;
cardiovascular disease; hypercholesterolasemia;
allele specific oligonucleotide; ss; probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human CETP gene allele specific oligonucleotide probe #54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       .9%; Score 13; DB 1; Length 15; 0.0%; Pred. No. 3.4e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 15 BP; 3 A; 8 C; 2 G; 2 T; 0 U; 0 Other;
                                                                                                                                                     Stephens JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 2.9%; Score 13; DB Best Local Similarity 100.0%; Pred. No. 3.4 Matches 13; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               oligonucleotide (ASO) primers that we polymorphisms in the human LCAT gene
                                                                                                                                                     Nandabalan K,
                                                                                                                                                                                                                                                                                                                                                  Claim 16; Page 17; 72pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADG98425/c
ID ADG98425 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               284 CACCCACTGCCAC 296
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                                                                                                                                                                                            WPI; 2002-557737/59.
                                                                                                                                                   Denton RR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO2003091277-A2
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                                                                                                                                                                  The invention comprises the amino acid and coding sequences of the human cholesteryl ester transfer protein (CETP), the invention also comprises polymorphisms identified within the CETP gene. The DNA and protein sequences of the invention are useful in haplotyping and/or genotyping the CETP gene in an individual. The DNA and protein sequences may also be used to screen drugs or compounds targeting the CETP or its variant to treat a condition or disease associated with CETP (e.g. atherosclerosis, eardiovascular disease or hypercholesterolaemia). The present DNA sequence represents an allele specific oligonucleotide probe for the
                                                                                                    New isolated polynucleotide useful for haplotyping and/or genotyping cholesteryl ester transfer protein (CETP) gene in an individual or in screening for drugs useful in treating diseases associated with CETP
                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Porcine reproductive and respiratory syndrome virus; PRRSV; vaccine; antigen; polymerase chain reaction; PCR; primer; ss.
                                                          Parks KE;
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0
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                                                          Lee HH,
                                                                                                                                                                                                                                                                         Sequence 15 BP; 4 A; 4 C; 5 G; 1 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lum MA;
                                                        Lachowicz M,
Tirrell C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Morozov I,
                                                                                                                                                 Claim 43; SEQ ID NO 57; 250pp; English.
                                                        Kazemi A, L
Sausker EA,
                                     (GENA-) GENAISSANCE PHARM INC
28-APR-2003; 2003WO-US013288.
                  26-APR-2002; 2002US-0375791P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRRSV sequencing primer DP966.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Halbur P,
                                                                                                                                                                                                                                                                                                                                 310 GCTCTGTCAGCCGCG 324
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               94US-00301435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              95WO-US010904
                                                                                                                                                                                                                                                                                                                                                                                                AAT14404 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                            15 GCTCTGTYAGCCTCG
                                                                                                                                                                                                                                                                                                               13; Conservative
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                                                       i, Chew i
                                                                                   WPI; 2003-865576/80.
                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
Matches 13; Conserv
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MENG X.
HALBUR P.
MOROZOV I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Paul PS, Meng X,
                                                                                                                                                                                                                                                         human CETP gene.
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                                                        Anastasio AE,
                                                                 Petersen N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-SEP-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO9606619-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-MAR-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                activity
                                                                                                                                                                                                                                                                                                                                                                                                                   AAT14404;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (PAUL/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (HALB/)
(MORO/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (LUMM/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MENG/)
                                                                                                                                                                                                                                                                                                                                                                               RESULT 410
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                                                                                                                                                                    Primer DP966 (AAT14404) is specific to porcine reproductive and respiratory syndrome virus (PRRSV). It was used with other sequencing primers (AAT14391-82) to determine the sequences of the putative membrane (M) and nucleocapsid (N) genes of PRRSV isolate ISU-12 (see also AAT14391-92) and of 5 other American PRRSV isolates (AAT14405-09) and Buropean
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
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New porcine reproductive and respiratory syndrome virus DNA - and proteins encoded by open reading frames of an Iowa strain of the virus; are used in vaccines against PRRSV in pigs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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0
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                                                                                                                                                                                                                                                                                                                                                                                    Sequence 16 BP; 2 A; 4 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17 BP; 5 A; 9 C; 2 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                            2.9%; Score 13; DB 100.0%; Pred. No. 3.6 tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mcswiggen J;
                                                                                                                     Page 77; 228pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 37; Page 56; 164pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hammerhead ribozyme substrate #4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99US-0129390P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAF01709 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     268 GGGCTTCTCCGG 280
                                                                                                                                                                                                                                                                                                                               strain Lelystad (AAT14410)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4 GGGCTTCTCCGG 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ribozyme; erythropoie
interferon alpha; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2000-647423/62.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200061729-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sapiens.
                                                                                                                  Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAF01709
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ò g ABZ62076;

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The sequences given in AAT80211-382 represent synthetic oligonucleotides which are complementary to a portion of the 5' untranslated region (UTR) of hepatitis C virus (HCV). These sequences may be used in a pharmaceutical composition for the control or prevention of HCV infection. They may be used to inhibit replication or expression of HCV in a sample. They may be used to inhibit replication or expression of HCV inhibit HCV replication in a cell and are therefor useful in the inhibit HCV replication in a cell and are therefor useful in the hepatocarcinoma. This sequence forms multiplex binding complexes with regions of the HCV genome. This sequence forms a duplex at the region - 227 and forms a purine strand triplex at the region -222 to -222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              go:nucleotide(s) complementary to HCV 5' untranslated region - used i treatment and detection of HCV infection, esp. hepatitis and hepato-
                                                                                                Complementary; 5' untranslated region; UTR; hepatitis C virus; HCV; inhibition; replication; expression; detection; chronic hepatitis; acute hepatitis; hepatocarcinoma; ss.
                                                                                                                                                                                                                                                                                                                              /*tag=' b
/note= "Comprises phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2.8%; Score 12.8; DB 1; Length 16; 31.2%; Pred. No. 3.8e+02; ve 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Reverse primer #142 used in multiplexing PCR/SBE assay.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kilkuskie RE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Seguence 16 BP; 0 A; 7 C; 6 G; 2 T; 1 U; 0 Other;
                                                          Oligo HCV-222, multiplex forming oligomer.
                                                                                                                                                                                                                                               1. .11
/*tag= a
/note= "2'-OMe RNA"
12. .16
/*tag= b
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hamlin HA,
Wolfe JL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (HOFF ) HOFFMANN LA ROCHE & CO AG F. (HYBR-) HYBRIDON INC.
                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 28; Page 22; 100pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   199 CCCTCCCGGGGACCTG 214
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               96WO-EP002427
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      95US-00471968
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CCCUCCGGGGGTCCTG 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       81.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAC73638 standard; DNA; 16
                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Goodchild J,
, Walther DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1997-043122/04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Oligo:nucleotide(s)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
Matches 13; Conserv
                                                                                                                                                                                                                             Key
modified_base
                                                                                                                                                                                                                                                                                                           modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-JUN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                             WO9639500-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Frank BL, G
Roberts PC,
                   16-OCT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                      12-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 02-FEB-2001
                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         carcinoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAC73638;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0 x 2 x 5 x 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ઠ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule, that modulates human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-theumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences ABZ66530 - ABZ66585 represent substrate/target sequences for the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a novel short interfering RNA (siRNA) nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                              Human, ribozyme; short interfering RNA, siRNA, HER2; K-Ras,
enzymatic nucleic acid, H-Ras, N-Ras, HIV, cytostatic, anti-HIV,
anti-rheumatic, cancer, AIDS, 8s.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2.9%; Score 13; DB 1; Length 17; 100.0%; Pred. No. 3.9e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Seguence 17 BP; 1 A; 4 C; 10 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 58; Page 129; 185pp; English
                                                                                                                                                                                                                                                                                        Human H-Ras DNAzyme target #867.
                                                                                                                                                               ABZ62076 standard; RNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAT80369 standard; DNA; 16 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29-MAY-2001; 2001US-0294140P.
06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-MAY-2002; 2002WO-US016840.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   225 CCTGCCCAGCCCC 237
                                                                                                                                                                                                                                                 (first entry)
                 2 GGTTGCGGAGGGT 14
                                           16 GGTTGCGGAGGGT 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 100.
Matches 13, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mcswiggen J;
                                                                                                                                                                                                                                               21-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-DEC-2002.
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RESULT 41
ABZ62076/
XX
ABZ62076/
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ABZ62076/
XX
ABZ62076/
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ABZ62076/
ABZ6
XX
ABZ62076/
ABZ6

Roberts NA;

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Gaps

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13

ठ 셤 RESULT 413

AAT80369

AAT80369 ID AAT8 XX AC AAT8

Query Match

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ABT34275/
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                                                                                                                                                                                                                                                                                                    The present invention relates to an oligonucleotide array comprising oligonucleotide tags fixed to a solid substrate. The oligonucleotide array is useful for genotyping a nucleic acid sample at one or more loci via single base extension (SBE) reactions. A pair of primers is used to amplify a polymorphic locus in a sample e.g. a single nucleotide bolymorphism (SMP). The present sequence is one of the primers used in the method of the present invention to amplify a polymorphic sample. The amplified nucleic acid product is then used as a template in a SBE reaction with an extension primer. The SBE reaction products are used to
          Oligonucleotide array; genotyping; single base extension reaction; SBE; PCR primer; polymorphic locus; single nucleotide polymorphism; ss.
                                                                                                                                                                                                                                         Universal array of oligonucleotides tags attached to a solid substrate along with locus-specific tagged oligonucleotides useful in genotyping using single base extension reactions.
                                                                                                                                                                                          Д;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                         Kaplan P, Lander ES, Lockhart
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hepatitis C virus; HCV; hepatocyte infection; non-A hepatitis;
non-B hepatitis; acute hepatitis; chronic hepatitis;
hepatocellular carcinoma; virucide; cytostatic; antisense therapy;
gene therapy; ss; DNA-RNA hybrid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Inhibitory oligonucleotide specific for hepatitis C virus #159
                                                                                                                                                                                                                                                                                                                                                                                                                                           2.8%; Score 12.8; DB 1; Length 16; 37.5%; Pred. No. 3.8e+02; very 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 16 BP; 2 A; 2 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                         (WHED ) WHITEHEAD INST BIOMEDICAL RES
                                                                                                                                                                                                                                                                                  Example 7; Page 63; 70pp; English.
                                                                                                                                                                                         Huang X,
                                                                                                                                                                                                                                                                                                                                                                                                     form the oligonucleotide array
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ВР
                                                                                                      27-MAR-2000; 2000WO-US008069.
                                                                                                                          99US-0126473P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             97US-00887505.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24 AGGGGTGGTGGCCATT 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AGGTTGGTGGCCAGT 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                     87.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABS65953 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                    (AFFY-) AFFYMETRIX INC
                                                                                                                                                                                        Hirschhorn JN,
                                                                                                                                                                                                                      WPI; 2000-656171/63
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                  Ryder T, Sklar P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US2002081577-A1
                                                              WO200058516-A2
                                          Unidentified.
                                                                                                                          26-MAR-1999;
23-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             02-JUL-1997;
                                                                                 05-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                14;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABS65953;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                        Fan J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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The invention describes synthetic oligonucleotides complementary to a portion of the 5' untranslated region of hepatitis C virus. The oligonucleotides may be used in methods for controlling, preventing, and treating hepatitis C virus infection, in antisense technology and gene therapy, and of detecting the presence of hepatitis C virus in a sample. Hepatitis C virus infects on moveloped, positive sense, single-stranded RNA virus which infects hepatocytes. HCV is the major cause of non-A, non-B, acute and chronic hepaticits, and has been associated with hepatocellular carcinoma. The invention describes methods and kits for inhibiting replication of HCV, inhibiting the expression of HCV nucleic acid and protein, and for treating HCV infections. This sequence represents a synthetic DNA-RNA hybrid oligonucleotide used for inhibiting HCV replication and expression of HCV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic oligonucleotides complementary to a portion of the 5' untranslated region of hepatitis C virus (HCV), useful for diagnosing and treating HCV infections and hepatocellular carcinoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eating disorder; polymorphism; dataset; allele; HGBASE identification; serotonin receptor 1D; delta-opioid receptor; dopamine receptor D2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .;
0
                                                                                                                                                                                                                                                                                                                                           Roberts PC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2.8%; Score 12.8; DB 1; Length 16; 81.2%; Pred. No. 3.8e+02; tive 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                           Wolfe JL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 16 BP; 0 A; 7 C; 6 G; 2 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   anorexia nervoŝa; bulimia nervosa; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Serotonin receptor 1D probe SEQ ID No 61.
                                                                                                                                                                                                                                                                                                                                        Goodchild J,
Walther DM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 31; Page 7; 74pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         214
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP.
95US-00471968.
96US-0021104P.
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20-JUL-2001; 2001US-0306440P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABT34275 standard; DNA; 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 2.8
Best Local Similarity 81.2
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                              Kilkuskie RL, Frau. -
'- 4A, Roberts NA,
                                                                                  KILKUSKIE R L.
                                                                                                         FRANK B L.
GOODCHILD J.
WOLFE J L.
ROBERTS P C.
HAMLIN H A.
ROBERTS N A.
                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-537132/57.
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06-JUN-1995;
02-JUL-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unidentified
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                                                                                  (KILK/)
                                                                                                                                                                                              (ROBE/)
(HAML/)
(ROBE/)
                                                                                                                                                                                                                                                                                 (WALT/)
                                                                                                                                       (GOOD/)
(WOLF/)
                                                                                                               FRAN/
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Huntington's disease.
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modified_base
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01-NOV-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                               invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAT12444;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                         The invention relates to a novel isolated nucleic acid molecule comprising a variant gene associated with an eating disorder and selected from any of 119 polymorphisms with their corresponding genotyphism in dataset, alleles and HGBASE identification, given in the specification. The novel nucleic acid molecule has polymorphisms in the serotonin receptor 1D, delta-opioid receptor, or dopamine receptor D2, which is useful in disgnostic and prognostic, assays for eating disorders, in particular anorexia nervosa and bulimia nervosa. This polymolectic seaquence represents a serotonin receptor 1D probe of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New substantially pure ribonucleic acid (RNA) complex comprising a first strand and a second strand that hybridize to each other, useful for treating cancer, systemic lupus erythematosus, Alzheimer's disease or
                                                                                                                                                                                                               New nucleic acid molecule having polymorphisms in the serotonin receptor 1D, delta-opioid receptor, or dopamine receptor D2, useful in diagnostic and prognostic assays for eating disorders, such as anorexia and bulimia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         88; RNA complex; immunosuppressive; cytostatic; cancer; systemic lupus erythematosus; Alzheimer's; Huntington's disease; salivary gland carcinoma; melanoma; brain tumour; leukaemia; lymphoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.8%; Score 12.8; DB 1; Length 16; 87.5%; Pred. No. 3.8e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 16 BP; 4 A; 10 C; 2 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mccallus DE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pyrimidine nucleotide flanking sequence 3.
                                                                                                                                                                                                                                                                                                                        Example 3; Page 60; 149pp; English
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  13-NOV-2001; 2001US-0331285P.
19-DEC-2001; 2001US-0340843P.
19-DEC-2001; 2001US-0340844P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18-OCT-2002; 2002US-0419532P.
28-OCT-2002; 2002US-0421757P.
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les 14; Conservative
                                                                                    (PRIC-) PRICE FOUND LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (NUCL-) NUCLEONICS INC.
                                                                                                                               Bergen AW, Yeager M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-348454/32.
                                                                                                                                                                       WPI; 2003-268122/26.
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                                                                                                                                                                                                                                                                                  nervosa
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Matches
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This invention relates to double stranded ribonucleic acid (RNA)

Structures and constructs. Specifically, it comprises first and second

RNA strands that hybridise to each other under physiological conditions

to form a double-strand region, wherein the double-strand region contains

cone or more mismatched regions that result in two or more double-stranded

segments. Furthermore, the mismatched regions may be cleaved by single-

strand ribonuclease enzymes. The present invention describes expression

vectors that encode dsRNAs with an intron containing exemplary target

cetrapy, these RNA complexes exhibit immunosuppressive and cytostatic

cativities and can be used to treat cancer, systemic lupus erythematosus,

Alzheimer's and Huntington's disease. The cancer is selected from,

melanoma, brain tumour, leukaemia and lymphoma. This oligonuclectide

sequence is a DNA flanking sequence used in an exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antiviral, phosphorothioate, mRNA 4; mRNA 5; herpes simplex virus 1; HSV; viral infection; HIV; varicella zoster virus; VZV; therapy; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antiviral phosphoro:thioate oligo:nucleotide(s) - active against e.g. herpes simplex virus 1, HIV and varicella zoster virus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ouery Match 2.8%; Score 12.8; DB 1; Length 16; Best Local Similarity 87.5%; Pred. No. 3.8e+02;
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/note= "phosphorothioate oligonucleotides"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indele
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Seguence 16 BP; 0 A; 10 C; 0 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Antiviral phosphorothioate oligonucleotide #27
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Example 9; Page 119; 204pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
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94JP-00268603
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(KAKE ) KAKEN PHARM CO LTD.
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12-JUL-1996;
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Young SA,
                                                                                                                AAX62953;
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                                                                    RESULT 420
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Matches
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                                                                                                                                                                                                                                                                                                                                    Antiviral; phosphorothioate; mRNA 4; mRNA 5; herpes simplex virus 1; HSV; viral infection; HIV; varicella zoster virus; VZV; therapy; ss.
         antiviral activity. These sequences, and the phosphorothicate oligonucleotides represented by AAT12418-T12434 (which are complementary to regions of the mRNA 4 or 5 of herpes simplex virus 1 (HSV)), are effective in the prevention and treatment of viral infection. The sequences are especially effective against infection by HSV, HIV or varicella zoster virus (VZV)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antiviral activity. These sequences, and the phosphorothioate oligonuclectides represented by AAT12418-T12434 (which are complementary to regions of the mRNA or 5 of herpes simplex virus 1 (HSV), are effective in the prevention and treatment of viral infection. The sequences are especially effective against infection by HSV, HIV or
                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antiviral phosphoro:thioate oligo:nucleotide(s) - active against e.g. herpes simplex virus 1, HIV and varicella zoster virus.
 AAT12435-T12454 represent phosphorothioate oligonucleotides with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAT12435-T12454 represent phosphorothioate oligonucleotides with
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                                                                                                              2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 4e+02; Live 0; Mismatches 2; Indels
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ilarity 87.5%; Pred. No. 4e+02;
Conservative 0; Mismatches 2; Indels
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                                                                                          Sequence 17 BP; 0 A; 2 C; 15 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 17 BP; 0 A; 3 C; 14 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                              Antiviral phosphorothioate oligonucleotide #26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Iwatani W,
                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Shimada J, Mizushima Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 6; Page 150; 163pp; Japanese.
                                                                                                                                                                                                                                             ВР
                                                                                                                                                             333 GGGGCGAGGCGAGG 348
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   varicella zoster virus (VZV)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (LTTL-) LTT INST CO LTD. (KAKE ) KAKEN PHARM CO LTD.
                                                                                                                                                                                                                                            AAT12443 standard; DNA; 17
                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                           Local Similarity 87.5
nes 14; Conservative
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Matches 14; Conserv
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modified_base
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                                                                                                                                                                                                                                                                                        17-SEP-1996
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                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                   AAT12443;
                                                                                                                Query Match
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                                                                                                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mcswiggen JA, Merlo PAO, Guo L, Skokut TA;
Merlo DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Maize, corn, Zea mays, delta-9 desaturase, GBSS, target, substrate, granule bound starch synthase, hammerhead ribozyme, hairpin ribozyme, modulation, gene expression; transgenic plant; cleavage, canola plant; caffeine synthesis, coffee plant; nicotine production; tobacco; fruit ripening; flower pigmentation; lignin production; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 expression - preferably modulates granule bound starch synthase in
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Delta-9 desaturase hamerhead ribozyme target SEQ ID NO:828.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ribozyme which modulates plant gene expression of DELTA-9 desaturase or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 38; Page 86; 155pp; English.
332 CGGGGGCGAGGCGAG 347
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nes 10; Conservative
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Folkerts O,
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Leukocyte Antigen (HLA)-DRB sequences. The primers are used in the methods of the invention. The specification describes a method for determining the HLA class II group type of a subject. The method for determining the HLA class II group type of a subject. The method comprises amplifying a target DNA sample from a subject using a particular HLA group-specific primer pair and determining whether a nucleic acid product is produced, therefore identifying the group type. Methods for determining the HLA class allele type of a subject are also described, where a specific HLA group-specific exon region primer pair is used. The methods are useful for determining the HLA class II type of a claim sample, by identifying the specific alleles present and determining the group specificity of alleles. The methods are diagnostically useful for histocompatibility analysis to see if donor and creipient groups match, and for further compatibility analysis
                                                                                                                                                                                                                                              Determining the Human Leukocyte Antigen Class II type Histocompatibility antigens - by using new intron-specific oligonucleotide primers for sequence specific primer PCR and sequencing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human leukocyte antigen class II type; HLA class II type;
histocompatibility locus antigen class II; PCR primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human leukocyte antigen class II type PCR primer DRB02.
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                                                                                                                                                                                                                                                                                                                            Claim 13; Fig 3A; 36pp; English
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                                                      97EP-00110438
                                                                                            97EP-00110438
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAX00304 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                          WPI; 1999-047888/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 14; Conserv
                                                                                                                                  (BIOT-) BIOTEST AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (BIOT-) BIOTEST AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26-JUN-1997;
                                                      26-JUN-1997;
                                                                                            26-JUN-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                  30-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EP892069-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Blasczyk R;
                                                                                                                                                                        Blasczyk R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAX00304;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 423
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            셤
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Detection of mutation(s) in the BRCAl gene - by hybridisation with an allele-specific oligo:nucleotide or by amplification, useful particularly for breast or ovarian cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAV20567-V20574 are probes used in a method to detect mutations in the ... human BRCA1 gene. Such mutations are used for classifying a tumour for diagnostic and prognostic purposes or detecting a predisposition of higher susceptibility to breast and ovarian cancer in an individual. The methods can be used for reducing the high incidence and mortality women at high risk. These women, once identified, can be targeted for more aggressive prevention programmes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human Leukocyte Antigen; HLA; HLA-DRB consensus sequence; intron 1; HLA class II group type; histocompatibility analysis; compatibility analysis; PCR primer; ss.
                                                                                                            Breast cancer; ovarian cancer; mutation; classification; detection; tumour; diagnostic; prognostic; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 4e+02; 2; Indels ive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Zeng
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Group-specific amplification primer for HLA-DRB02.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17 BP; 8 A; 1 C; 7 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Olson SJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 4; Page 40; 62pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    White MB,
                                                                                                                                                                                                                                                                                                                                            96US-0023184P.
96US-0023187P.
96US-002323P.
96US-0022421P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           366 GCAGGAAGAGGAACGG 381
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAX04779 standard; DNA; 17 BP
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                                                                                                                                                                                                                                                                                                            97WO-US013654
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                                    02-JUL-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 87.5
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Murphy PD, Allen AC,
                                                                          Human BRCAl probe #4.
                                                                                                                                                                                                                                                                                                                                                                                                                                              (ONCO-) ONCORMED INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1998-159166/14.
                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                    05-AUG-1996;
06-AUG-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-APR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                WO9805677-A1
                                                                                                                                                                                                                                                                                                            04-AUG-1997;
                                                                                                                                                                                                                                                                                                                                              05-AUG-1996;
05-AUG-1996;
                                                                                                                                                                                                                                                                      12-FEB-1998
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                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
AAV20570;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAX04779;
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AAX04779

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Gape

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2; Indels

DB 1; Length 17;

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Action has been developed of determining the Human Leukocyte Antigen class II (HLA Class II) group type of a subject. The method comprises:

(i) amplitying a target DNA sample from a subject using a particular HLA group-specific primer pair (sequence specific primer PCR - SSP-PCR); and group-specific primer pair (sequence specific primer PCR - SSP-PCR); and (ii) determining whether a nucleic acid product is produced, therefore identifying the group type. AAX00303 to AAX00396 represent specifically clasmed oligonucleotide primer for use in the above method. These oligonucleotides are useful for determining the HLA Class II type of a patient sample, by identifying the specific alleles present and cetermining the group specificity of alleles. Steps (i) and (ii) in the method are diagnostically useful for histocompatibility analysis to see if donor and recipient groups match. The new sequences are useful for providing an insight into the genetic relationship between different alleles of HLA Class II genes. The high resolution, nucleic acid based method using the intron-specific primers is more efficient than prior art methods using exon based primers, as few exon sequences offer conserved primer binding sites, resulting in a limited number of primer pairs and insufficient specificity for alleles, as allelic variations exist between the primer sites. The SSP-PCR method allows separation of haplotypes in 95% of patient samples, allowing resolution of cis-trans linkages of the preserved primer sites. The samples cannot be achieved with other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                               Determining the Human Leukocyte Antigen Class II type Histocompatibility antigens - by using new intron-specific oligonucleotide primers for sequence specific primer PCR and sequencing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antisense oligonucleotide, multiple target, antisense treatment; impaired respiration; inflammation; lung disease; pulmonary vasconstriction; inflammation; allergic rhinitis; acute asthma; allergy; asthma; impeded respiration; respiratory distress syndrome; pain; cystic fibrosis; pulmonary hypertension; pulmonary vasconstriction; emphysema; chronic obstructive pulmonary disease; leukemia; lymphoma; carcinoma; colon cancer; breast cancer; lung cancer; pancreatic cancer; hepatocellular carcinoma; kidney cancer; melanoma; hepatic metastasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17 BP; 2 A; 10 C; 4 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C/EBP-beta antisense oligonucleotide fragment.
                                                                                                                                                                                                                               Claim 4; Fig 3; 36pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            232 AGCCCCCCGAACCCCGC 247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAX55127 standard; DNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             98WO-US019419.
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98US-00093972.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 AGCGCCCCCCCCCCC 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity 87.5%;
nes 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
WPI; 1999-083585/08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  prostate cancer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-JUL-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-SEP-1997;
09-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAX55127;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 424
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAX55127,
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The specification describes antisense oligonucleotides (AAX52869-X55271)

directed against at least 2 mRNAs selected from target genes, coding and
non-coding regions of RNAs corresponding to target genes, gene initiation
codons, genomic flanking regions, intron-exon borders, the 5'-end, the 3'
codons, genomic flanking regions, intron-exon borders, the 5'-end, the 3'
codons, genomic flanking regions, intron-exon borders, the 6'-end, the 3'
codons, mixtures. The antisense oligonucleotides may be derived
conditions or mixtures. The antisense oligonucleotides may be derived
from sequences AAX55272-74. These multiple target oligonucleotides
conditions. Typical diseases and conditions are those
associated with impaired respiration and inflammation, including lung
diseases, pulmonary vasoconstriction, inflammation, allergic rhinitis,
acute asthma, allergies, asthma, impeded respiration, respiratory
diseases (COPD), and cancers such as leukemias, lymphomas, carcinomas e.g.
colon cancer, breast cancer, lung cancer, pannoral metastassized
colon cancer, breast cancer, lung cancer, panoreallular cancer, panters metastasized
colon cancer, breast cancer, lung cancer, panters metastasized
colon cancer, breast cancer, lung cancer, panters metastasized
colon cancer, breast cancer, lung cancer, panters metastasized
colon cancer which may metastasize or have metastasized
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               phosphorothioate; impaired respiration; inflammation; allergy; allergy; allergic disease; bronchoconstriction; inhibitor; antiinflammatory; antiallergic; antiasthmatic; cytostatic; analgesic; impaired airway; lung disease; ischaemic condition; pulmonary vasoconstriction; asthma; respiratory distress syndrome; pain; cystic fibrosis; emphysema; pulmonary hypertension; chronic obstructive pulmonary disease; COPD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                      New antisense oligonucleotides used in treatment of, e.g. pulmonary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; adenosine receptor; low adenosine antisense oligonucleotide;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cancer; leukaemia; lymphoma; carcinoma; metastasis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17 BP; 0 A; 12 C; 5 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                            Disclosure; Page 71; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   272
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                   (UYEC-) UNIV EAST CAROLINA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAA34574 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-JUL-2000 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16 geggeegeegeege
                                                                                                           WPI; 1999-229400/19
                                                                                                                                                                              vasoconstriction
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAA34574;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                Nyce JW;
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 425
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The present invention describes low admonstrated by conjugational the antisense oligonucleotides and compositions (I) comparising them. In the antisense oligonucleotides the A is replaced by a 'Universal' or alternative base. (I) can have respiratory, bronchodilator, antihinflammatory, analgeaf., immunosuppressive, antiaethmatic, hypotensive and cytostatic activities. The antisense oligonucleotides and (I) can be used to down-regulate the expression and or activity of target polypeptides associated with cump/respiratory disorders and malignacies, auch as stimulating and activating peptide factors and transmitters, transcription factors, immunoglobulins and antibodies, antibody receptors, cytokines and chemokines, endogenously produced specific and non-apecific canzymes, chemokines, adhesion molecules and their receptors, central nervous system (CNS) and peripheral nervous and non-nervous system ceceptors, defensins, growth factors, vasocative peptides and receptors, binding proteins and malignancy associated proteins. The receptors, binding proteins and malignancy associated proteins. The cansmitters defensins, growth factors, vasocative peptides and receptors, binding proteins and malignancy associated disorders including respiratory obstruction (especially pulmonary obstruction and/or bronchoconstriction) and/or lung inflammation, allergy(ies) and/or condition and formulance associated with a disease or condition and formulance associated with a disease or conditions.
                                                                                                                                                                                                                                                                                                                               Low adenosine (A) content antisense oligonucleotides which do not trigger adenosine receptors during metabolism, useful e.g. for treating cancers
                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 14; Page 265; 1592pp; English
                                                                                   24-MAR-2000; 2000WO-US008020
                                                                                                                                                                                                                                                                                                                                                                                and respiratory obstructions
                                                                                                                                                                         (UYEC-) UNIV EAST CAROLINA.
                                                                                                                                                                                                                                                                                     WPI; 2000-679539/66.
                                                                                                                                                                                                (NYCE/) NYCE J W.
WO200062736-A2
                                                                                                                                 06-APR-1999;
                                            26-OCT-2000
                                                                                                                                                                                                                                            Nyce JW;
  8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              g
                                                                                                                                                                                                                                                                            The present invention describes a new composition comprising an antisense oligonucleotide (ON) with low adenosine (up to 15%), which targets concletic acids involved in bronchoconscriction, allergies, and/or inflammation. The ON can have antinflammatory, antiallergic, antiasthmatic, cytostatic and analgesic activities. The compositions are useful for the treatment of diseases associated with inflammation, impaired airways, including lung disease associated with inflammation, impaired airways, including lung disease and diseases whose secondary effects afflict the lungs of a subject. They can be used for treating c. g. ischaemic conditions, pulmonary vasoconstriction, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, pulmonary hypertension, emphysema, chronic obstructive calcinomas, and cancers which may metastasise to the lungs, including breast and prostate cancer. The reduction of the adenosine content of the calcuses aide effects. The A-containing ONs break down with the release of deoxyadenosine which activates adenosine receptors causing bronchoconstriction and inflammation. AAA32313 to AAA35312 represent the invention, which correspond to SEQ ID No:1 to 185; but the sequences differ from the previously named sequences: SEQ ID No:1 to 1860 (AAA32331 to AAA3392) are specifically claimed ONs from the present invention. N.B. Sequences given in the disclosure of the present invention do not match is the interface sequence given in the sequences given in the sequences given in the sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Low adenosine antisense oligonucleotide; phosphorothioate; allergy; human; airway disorder; bronchoconstriction; lung inflammation; human; airway disorder; bronchoconstriction; lung inflammatory; immunosuppressive; antisthmatic; analgesic; hypotensive; cytostatic; respiratory obstruction; pulmonary obstruction; impeded respiration; surfactant hypoproduction; pulmonary vasoconstriction; asthma; RDS; respiratory distress syndrome; pani; cystic fibrosis; allergic rhinitis; pulmonary hypertension; emphysema; pulmonary transplantation rejection; chronic obstructive pulmonary disease; pulmonary infection; bronchitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                               New antisense oligonucleotides useful for treating e.g. pulmonary vasoconstruction, inflammation, allergies, asthma, hypertension, bronchitis, emphysema, respiratory distress syndrome, ischemia or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 4e+02; tive 0; Mismatches 2; Indel8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 17 BP; 0 A; 12 C; 5 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human C/EBP polynucleotide fragment #2263.
                                                                                                                                                                                                                                            Disclosure; Page 548; 1343pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAF20696 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               257 GCGGTCGGCCCGGGGC 272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16 GCGGGCGCGCGCGGC 1
  (UYEC-) UNIV EAST CAROLINA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14-MAR-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14; Conservative
                                                                                     WPI; 2000-205971/18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAF20696;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cancer;
                                              Nyce JW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                426
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              8
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present invention describes low adenosine (A) content antisense

99US-0127958P

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condition selected from pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome (RDS), pain, cystic fibrosis (CF), allergic rhinitis (AR), pulmonary hypertension, emphysema, chronic obstructive pulmonary disease (COPD), pulmonary transplantation rejection, pulmonary infections, bronchitis, and/or cancer. AAF18434 to AAF21543 represent human polynucleotide fragments and antisense oligonucleotides used in the exemplification of
                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ribozyme; erythropoietin; granulocyte colony stimulating factor;
                                                                                                                                                                                                                                                                                                                                     .
0
                                                                                                                                                                                                                                                                          2.8%; Score 12.8; DB 1; Length 17;
87.5%; Pred. No. 4e+02;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                Sequence 17 BP; 0 A; 12 C; 5 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hammerhead ribozyme substrate #2955.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAF06158 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16-FEB-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                       257 GCGGTCGGCCCGGGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 87.5
nes 14; Conservative
                                                                                                                                                                                   the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                      16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAF06158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 427
                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAF06158
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The invention relates to the construction of a vector which comprises a multiple number of Shear Stress Response Elements (SSRE) from various can gene promoter sequences and one or more genes, antisense molecules, cribozymes, double stranded RNA, or a nucleic acid which encodes a creativity of the protein or peptide. This sequence represents the SSRE sequence from the promoter of the platelet-derived growth factor A (PDGF-CA). The vector is useful for stimulating or inhibiting vascular endothelial cell or capillary endothelial cell proliferation and for stimulating angiogenesis in cells. The vector or gene of interest is useful for modulating vascular permeability in a mammal, for stimulating or inhibiting the formation, maturation or regression of blood vessels, modulating genes or proteins involved in a diseases, down regulating angiogenesis and for treating vasculogenic angiogenic disorders. These disorders include cardiovascular disorder, neoplastic disorders. These disorders include cardiovascular disorder, hypercholesterolaemia
                                                                                                              Expression vector comprising multiple shear stress response elements, useful for modulating endothelial cell proliferation, stimulating or down regulating angiogenesis and treating vasculogenic/angiogenic disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel nucleic acid molecule e.g., ribozymes or antisense nucleic acid molecules, which downregulates expression of a checkpoint kinase-1 gene, useful for treating colorectal, lung, breast or prostate cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     checkpoint kinase-1; Chk1; antisense; ribozyme; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mcswiggen J, Booher RN, Holman PS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ouery Match 2.8%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 40+02; Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 17 BP; 0 A; 2 C; 15 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human Chkl ribozyme substrate SEQ ID NO: 7.
                                                                                                                                                                                                           Example 1; Page 45; 61pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              333 GGGGCGAGGGCGAGG 348
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               03-FEB-2000; 2000US-0179983P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAH94582 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cancer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Jarvis T,
                                                                    WPI; 2000-452382/39.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-496922/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FATT/) FATTAEY A R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and wound healing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200157206-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RNA cleavage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Fattaey AR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-OCT-2001
                        Resnick N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAH94582;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 429
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAH94582/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, EAR3/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoietin, granulocyte colony stimulating factor protein and
                                                                                                                                                                                                                                                                                                                                                                                                                      Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cytostatic, cardiant, vasotropic, vulnerary, antidiabetic, hypotensive, antiatherosclerotic, antilipemic, gene therapy, vector, SSRE, promoter, Shear Stress Response Element, antisense, riboxyme, repressor antibody, platelet derived growth factor A, PDGF-A, angiogenesis, ischaemia, cardiovascular disorder, neoplastic disorder, atherosclerosis, ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hypertension; diabetes; hypercholesterolaemia; wound healing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2.8%; Score 12.8; DB 1; Length 17; 37.5%; Pred. No. 4e+02; rative 8; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17 BP; 2 A; 4 C; 1 G; 0 T; 10 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Shear Stress Response Element from PGDF-A gene.
                                                                                                                                                                                                                                                                                                                                Mcswiggen J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 42; Page 123; 164pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           97 TGTTTTTCTCGCTGAC 112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAA70569 standard; DNA; 17 BP
                                                                                                                                                                                       11-APR-2000; 2000WO-US009721.
                                                                                                                                                                                                                                                                                                                              Blatt L, Zwick M, Pavco P,
                                                                                                                                                                                                                                   99US-0129390P
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                                                                                                                                                                                                                                                                               (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
interferon alpha; ss
                                                                                                                                                                                                                                                                                                                                                                          WPI; 2000-647423/62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
les 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   interferon alpha
                                                                                          WO200061729-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200039275-A2.
                                                Homo sapiens.
                                                                                                                                                                                                                                      12-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
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                                                                                                                                      19-OCT-2000
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AAA70569

Query Match

Best Loc Matches

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Gaps

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Claim 4; Page 52; 115pp; English

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BP

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ABK00059 standard; RNA; 17
                               ABK00059;
ABK00059/
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        downregulating the expression of the human checkpoint kinase-1 (Chkl) the gene. These may be antisense or ribozyme sequences, and are useful in the treatment of diseases associated with conditions affected by Chkl levels, including cancer. The present sequence is an oligonucleotide described in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention provides nucleic acid molecules capable of downregulating the expression of the human checkpoint kinase-I (Chkl) gene. These may be antisense or ribozyme sequences, and are useful in the treatment of diseases associated with conditions affected by Chkl levels,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            including cancer. The present sequence is an oligonucleotide described in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel nucleic acid molecule e.g., ribozymes or antisense nucleic acid molecules, which downregulates expression of a checkpoint kinase-1 geruseful for treating colorectal, lung, breast or prostate cancers.
                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                     Human; checkpoint kinase-1; Chk1; antisense; ribozyme; gene therapy;
The present invention provides nucleic acid molecules capable of
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Holman PS;
                                                                                          ch 2.8%; Score 12.8; DB 1; Length 17; I Similarity 87.5%; Pred. No. 4e+02; 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Booher RN,
                                                                       Sequence 17 BP; 5 A; 2 C; 7 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Seguence 17 BP; 5 A; 2 C; 7 G; 0 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                               Human Chkl ribozyme substrate SEQ ID NO: 959.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mcswiggen J,
                                                    the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 4; Page 77; 115pp; English
                                                                                                                                                                                                               AAH95534 standard; RNA; 17 BP
                                                                                                                                       272 CTTCTCCGGAGGCACC 287
                                                                                                                                                                                                                                                                                                                                                                                                   02-FEB-2001; 2001WO-US003504
                                                                                                                                                                                                                                                                                                                                                                                                                       03-FEB-2000; 2000US-0179983P.
                                                                                                                                                          crrcrccaraccacc 2
                                                                                                                                                                                                                                                                                                                                                                                                                                            RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                RNA cleavage; cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Jarvis T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                     FATTAEY A R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-496922/54.
                                                                                                                                                                                                                                                                                                                                                        WO200157206-A2.
                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                         09-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fattaey AR,
                                                                                                                                                                                                                                                                                                                                                                              09-AUG-2001.
                                                                                                                                                          17
                                                                                                                                                                                                                                    AAH95534;
                                                                                           Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                            (RIBO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                        (FATT/)
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The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down cregulates expression of a neutite growth inhibitor gene (NGCD). The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a DNAzyme) an Inozyme (an endolytic nucleic acid cleaving an RNA molecule possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN motif) property of the a YGY motif). The CD20-cargeting nucleic acid is used to cleave RNA with a YGY motif). The CD20-cargeting nucleic acid is used to cleave RNA cyller and the presence of a divalent cation that is preferably Mg^2+.

Furthermore, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more control to the coll and treat a patient having a condition associated with the level of the TD30 in the presenta, B-cell lymphoma, low-grade or follicular non-thody inmunocytoma (NHL), immunocytoma (NHL), immunocytoma (MCL), immunocytoma (MCL),
                                                                                                                                                                                                            Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme; DNAzyme; inczyme; dever; amberzyme; zinzyme; luxphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphomy; leukaemia; human immunodeficiency virus; HIV associated NHL; nantle-cell lymphoma; MCI; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arbropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; Parkinson's disease; ataxia; Huntington's disease; creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.
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                                                                                                                  Human NOGO Hammerhead Ribozyme #59.
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(CHOW/)
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Gaps

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2; Indels

0; Mismatches

272 CTTCTCCGGAGGCACC 287

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14; Conservative

Matches

17 criciccaradecace 2

Query Match 2.8%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 4e+02;

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presence of a divalent cation that is preferably Mg^2^+. Furthermore, the nucleic acid may be contacted with a cell to reduce NOGO activity of the cell and treat a patient having a condition associated with the level of NOGO. The treatment may further comprise the use of one or more therapies. In particular, the NOGO-targetting nucleic acid may be used to treat central nervous system (CMS) injury and cerebrovascular accident (CVA, stroke), Alzheimer's disease, dementia, multiple sclerosis (MS), chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (MS), darkinson's disease, ataxia, Huntington's disease, Creutzfeldt-Jakob disease, muscular dystrophy, and/or other neurodegenerative disease sequence is a hammerhead ribozyme of the invention
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MCSWIGGEN J.
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The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down control expression of a neurite growth inhibitor gene (NAGO). The nucleic acids (e.g. a ribozyme or a nuorate acids may be enzymatic nucleic acids (e.g. a ribozyme or a nuorate (an endolytic nucleic acid cleaving an RNA molecule DNAzyme) an Inozyme (an endolytic nucleic acid cleaving an RNA molecule DNAzyme) an Inozyme (an endolytic nucleic acid cleaving RNA with an NGN triplet), a zinzyme (cleaving RNA with a pationt having a condition associated with the level CC therapies. In particular, the CD20 targetting nucleic acid with the lavel CC treat lymphoma (NHL), bulky low-grade or follicular NHL, lymphocytic lymphoma (MCL), immunocytopaenia, B-cell lymphoma, low-grade or follicular nor-lowdexima (MCL), immunocytopaenia, and inflammatory arthropathy. The NOGO. The treatment may further comprise the use of one or more crell and treat a patient having a condition associated with the lavel of presence of a divalent cation that is preferably Mg^2+. Furthermore, the nucleic acid may be contacted with a cell to reduce NOGO activity of the contacted with a cell to reduce NOGO activity of the cell and treat a patient having a condition associated with the lavel of therapies. In particular, the NOGO-targetting nucleic acid may be used to creat central nervous system (CNA, stroke), Alzheimer's disease, dementia, multiple sclerosis (MS), chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (MS), chemotherapy-induced neuropathy, amyotrophic acide may reduce of these miscular minodace 
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                                          Claim 88; Page 66; 200pp; English.
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WO200159103-A2

Synthetic

Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOG0; hammerhead ribozyme; DNAzyme; inozyme; derla yrowth inhibitor gene; NOG0; hammerhead ribozyme; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; NHL; lymphocytic leukaemia; Mnman immunodeficiency virus; HIV associated NHL; mantle-cell lymphoma; MCL; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; Parkinson's disease; ataxia; Huntington's disease; creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.

Human NOGO Hammerhead Ribozyme #58.

12-MAR-2002 (first entry)

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The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down capulates expression of a neurite growth inhibitor gene (NGO). The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a DNAzyme) an Inozyme (an endolytic nucleic acid cleaving a an RNA molecule DNAzyme) an Inozyme (an endolytic nucleic acid cleaving a an RNA molecule CC possessing an NCH mocif), a G-leaver (cleaving RNA with a NCH worlf). The CD20-targetting nucleic acid is used to cleave RNA with a YGY motif). The CD20-targetting nucleic acid is used to cleave RNA cc f CD20 in the presence of a divalent cation that is preferably MG<sup>2</sup>+. The CD20 targetting nucleic acid is used to cleave RNA cc f CD20. The treatment may further comprise the use of one or more therapies. In particular, the CD20 targetting nucleic acid may be used to treat lymphoma, leukaemia, HG-CD30 targetting nucleic acid may be used to treat lymphoma (MLI), bulky low-grade or follicular non-Hodgkin's lymphoma (MLI), immunocytoma (IMC), small B-cell lymphocytic lymphoma, immunodeficiency virus) associated NHL, mantle-cell lymphoma (MCI), immunocytoma (IMC), small B-cell lymphocytic lymphoma, immunodeficiency virus) associated NHL, mantle-cell cr argetting nucleic acid is used to cleave RNA of the NOGO activity of the consence of a divalent cation that is preferably MG<sup>2</sup>2+. Furthermore, the nucleic acid may be contacted with a cell to reduce NOGO activity of the cell and treat a partient having a condition associated with the level of the reatment may further comprise the use of one or more therapies. In particular, the NOGO-targetting nucleic acid may be used to the central nervous system (CNS) injury and cerebrovascular accident (CVM, stroke), Alzheimer's disease, dementia, multiple aclerosis (ALS), chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (ALS), disease, muscular chica enervo en enerver neuropathy, and/or the enervolation enervolated enervolated energy energetting neuropat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chowrira BM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 88; Page 98; 200pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                central nervous system injury.
                                                                                                 09-FEB-2001; 2001WO-US004273.
                                                                                                                                                                                                       2000US-0181797P
                                                                                                                                                                                                                                                             28-FEB-2000; 2000US-0185516P.
06-MAR-2000; 2000US-0187128P.
                                                                                                                                                                                                                                                                                                                                                                                                               (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Blatt L, Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MCSWIGGEN J.
CHOWRIRA B M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-607195/69.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BLATT L.
                                                                                                                                                                                                       11-FEB-2000;
16-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                               (BLAT/)
(MCSW/)
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11-FEB-2000; 2000US-0181797P. 28-FEB-2000; 2000US-0185516P. 06-MAR-2000; 2000US-0187128P.

(RIBO-) RIBOZYME PHARM INC.

BLATT L.

(MCSW/) (BLAT/)

09-FEB-2001; 2001WO-US004273

WO200159103-A2.

16-AUG-2001

sapiens.

Ношо

Synthetic.

Chowrira BM;

Blatt L, Mcswiggen J, CHOWRIRA B M MCSWIGGEN J.

WPI; 2001-607195/69.

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The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down cregulates expression of a neutite growth inhibitor gene (NDGO). The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a DNAzyme) an Inozyme (an endolytic nucleic acids (e.g. a ribozyme or a maberzyme). The CD20-targetting nucleic acid is used to cleave RNA with a NYN mouth; prometry a resence of a divalent cation that is preferably Mg<sup>2</sup>+. The CD20-targetting nucleic acid is used to cleave RNA with a YGY motif). The CD20-targetting nucleic acid is used to cleave RNA with a NGN triplet), a zinzyme (cleaving RNA with a PG20 in the presence of a divalent cation that is preferably Mg<sup>2</sup>+. The cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the use of one or more therapies. In particular, the CD20 targetting nucleic acid may be used to treat lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular non-Hodgkin's lymphoma (NHL), bulky low-grade or follicular NHL, lymphocytic lymphoma, leukaemia, B-cell lymphoma, low-grade or follicular NHL, lymphocytic lymphoma (MCL), immunocytopaenia, and inflammatory arthropathy. The NOGO-treatment may further comprise the used of the NOGO gene in the presence of a divalent cation that is preferably Mg<sup>2</sup>2+. Furthermore, the nucleic acid may be contacted with a cell to reduce NOGO activity of the capter a patient having a condition associated with the level of NOGO. The treatment may further comprise the use of one or more therapies. In particular, the NOGO-targetting nucleic acid may be used to cleave RNA of ene of one or more therapies. In particular, the NOGO-targetting nucleic acid may be used to cleave the contact a patient having a condition associated with the level of the contact and may further comprise the used of one or more treatment and result and result may further comprise the used of one or more treatment and result and result and result and result 
Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (CVA, stroke), Alzheimer's disease, dementia, multiple sclerosis (MS),
                                                                                                                                                                                                                                                                                                                                                                                          Claim 88; Page 66; 200pp; English.
                                                                                                                                                                                                                                    central nervous system injury.
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Gaps

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2; Indels

0; Mismatches

254 GCCGCGGTCGGCCCGG 269

14; Conservative

Matches

17 GCCGCGGACAGCCCGG

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ABK00058 standard; RNA; 17 BP

RESULT 434

ABK00058

Query Match 2.8%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 4e+02;

BP

ABV85745 standard; DNA; 17

ABV85745/

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to oligonucleotides that downregulate the expression of human Grb2-related with Insert Domain (GRID) gene. GRID is a T-cell co-stimulatory adaptor protein. The oligonucleotides are useful for modulating the expression of GRID, to treat conditions such as tissue/graft rejection and leukaemia. The oligonucleotides can also be administered in conjunction with other therapies such as radiation, chemotherapy and cyclosporin treatment. The present oligonucleotide was
     Chemocherapy-induced neuropathy, amyorrophic lateral sclerosis (ALS), Parkinson's disease, ataxia, Huntington's disease, returifiedtalakob disease, muscular dystrophy, and/or other neurodegenerative disease states which respond to the modulation of NOGO expression. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Domain
                                                                                                                                                                              Gaps
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co-stimulatory adaptor protein; tissue rejection; graft rejection;
leukaemia; cytostatic; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleic acid(s) for regulating the Grb2-related with Insert [ID] gene comprises using antisense and enzymatic nucleic acid
                                                                                                                                                                             ·;
                                                                                                                                       cch 2.8%; Score 12.8; DB 1; Length 17; al Similarity 87.5%; Pred. No. 4e+02; 14; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                Human GRID NCH ribozyme substrate oligonucleotide #152.
 chemotherapy-induced neuropathy, amyotrophic lateral
                                                                  sequence is a hammerhead ribozyme of the invention
                                                                                                    Sequence 17 BP; 0 A; 6 C; 3 G; 0 T; 8 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17 BP; 3 A; 4 C; 9 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mcswiggen JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  molecules such as hammerhead ribozymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 4; Page 65; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          the invention
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                                                                                                                                                                                                                                                                                                                             ABL46698 standard; RNA; 17
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                               ABL46698;
                                                                                                                                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes an isolated nucleic acid (I) encoding a human UDP-GalNAc:polypeptide N-acetylgalactosaminyltransferase 10 (pp-GaNTaee 10, EC 2.4.1.41) protein. Human pp-GaNTaee 10 is located to chromosome 7411.2. (I) can be used in gene therapy. Molecules of the present invention can be used in therapy, particularly to prevent or treat a disorder associated with decreased expression or activity of pp-GaNTaes. The sequences given in ABV85011 to ABV86689 and ABP5150 to The sequence data for this patent is not represented in the printed present incention. N.B. The sequence data for this patent is not represented in the printed process.
                                                                                                   Human; UDP-GalNAc:polypeptide N-acetylgalactosaminyltransferase 10;
pp-GaNTase 10; EC 2.4.1.41; chromosome 7q11.2; gene therapy; scanning;
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                                                                         Human pp-GaNTase 10 scanning 17-mer SEQ ID NO:738
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30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000670.
23-MAX-2001; 2001US-00864761.
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2001WO-US000666
2001WO-US000667
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                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       encoded protein.
                                                                                                                                                               sapiens.
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30-JAN-2001;
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Local Sir.
14;
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                                                                                                                                                                             Synthetic.
                ABV85745;
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RESULT 437 ABV85746/c

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200 CCTCCCGGGGACCTGC 215

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16 ccrcccrccciccic 1

87.58;

14; Conservative

Matches

Local Similarity

Query Match

(first entry)

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Chromosomal genomic alteration; genome altering oligonucleotide; PCR; 88; o-methyl modification; LNA modification; phosphorothioate linkage; bNA repair; DNA alteration; environmental tolerance; hygromycln-B; abiotic stress tolerance; improved nutritional value; hygromycln: Brimer; amino acid over production; herbicide resistance; glyphosate resistance; porphyric herbicide resistance; sulphonylurea herbicide resistance; porphyric herbicide resistance; sulphonylurea herbicide resistance; modified oil production; modified starch production; waxy starch; modified foral morphology; male-sterile plant; albino musant; modified fatty acid content; reduced plant; albino musant; increased stearate production; albino plant; increased stearate production; albino plant;
                                                    Male-sterile plant producing genome altering oligonucleotide #291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-JUN-2001; 2001WO-US017672.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-JUN-2000; 2000US-0208538P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-OCT-2000; 2000US-0244989P-27-MAR-2001; 2001US-00818875
                                                                                                                                                                                                                                                                                                    photosynthetic process.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (UYDE ) UNIV DELAWARE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-106307/14.
                                                                                                                                                                                                                                                                                                                                                                                                WO200192512-A2
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                 09-APR-2002
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                                                                                                                                                                                                                                                                                                                                                         Synthetic.
Nucleic acid encoding human UDP-GalNAc:polypeptide N-cetylgalactosaminyltransferase 10 protein is useful to diagnose, prevent and treat disorders associated with reduced or over expression of the encoded protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     human UDP-GalNAc:polypeptide N-acetylgalactosaminyltransferase 10 (pg-GaNTase 10, EC 2.4.1.41) protein. Human pp-GaNTase 10 is located to chromosome 7q11.2. (i) can be used in gene therapy. Molecules of the present invention can be used in therapy, particularly to prevent or treat a disorder associated with decreased expression or activity of pp-GaNTase. The sequences given in ABV85011 to ABV86699 and ABP53502 to ABP53504 are given in the exemplification of the present invention. N.B. The sequence data for this patent is not represented in the printed specification but is based on sequence information supplied by the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes an isolated nucleic acid (I) encoding a
                                                                                                                                             Human, UDP-GalNAc:polypeptide N-acetylgalactosaminyltransferase 10;
pp-GaNTase 10; EC 2.4.1.41; chromosome 7q11.2; gene therapy; scanning;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.8%; Score 12.8; DB 1; Length 17;
87.5%; Pred. No. 4e+02;
ative 0; Mismatches 2; Indels
                                                                                                           Human pp-GaNTase 10 scanning 17-mer SEQ ID NO:739.
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2001WO-US000670.
2001US-00864761.
2001US-0315984P.
ABV85746 standard; DNA; 17 BP
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2001WO-US000664.
2001WO-US000665.
2001WO-US000666.
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                                                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (AEOM-) AEOMICA INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-724954/79.
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30-JAN-2001; 2
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                                                                                                                                                                                                                                             Synthetic.
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                                     ABV85746
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Kim J;

Rice MC,

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The invention relates to an oligonucleotide for targeted alteration of a genetic sequence, which comprises a single-stranded oligonucleotide having a DNA domain. The DNA domain has at least one mismatch with respect to the genetic sequence to be altered and further comprises chemical modifications of the oligonucleotide. The chemical modifications of the oligonucleotide. The chemical modifications of the oligonucleotide are useful for more of these modifications. The oligonucleotides are useful for more of these modifications. The oligonucleotides are useful for coligonucleotides are particularly useful for creating plants with desired phenotypes, e.g. environmental or abjoic stress tolerance, improved conferring amino acid over production), herbicide resistance (e.g. altering amino acid content of plants or conferring amino acid over production), herbicide resistance (e.g. altering amino acid subhonylures herbicide resistance), disease resistance, modified oil production, modified starch production (e.g. increased starch or production of waxy starch), altered floral morphology (e.g. male-sterile plants) or modified fatty acid content (e.g. increased starch or production of waxy starch), altered floral morphology (e.g. male-sterile plants) or modified starty acid content (e.g. increased starch increased stearte or reduced linolenic acid). The oligonucleotides are also useful for producting albino mutants for the analysis of photosynthetic processes. This sequence represents a genome
New oligonucleotides with modified nuclease-resistant termini, useful for creating plants with desired phenotypes, e.g. stress tolerance, improved nutritional value, herbicide or disease resistance, or modified oil
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                                                                                                                                                                                                                                                                                                                         Claim 7; Page 87; 220pp; English.
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Gaps

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2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 4e+02; tive 0; Mismatches 2; Indels

Conservative

Best Local Similarity Matches 14; Conserv

Query Match

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Gaps

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343 GCGAGGTTCAGGCCTT 358

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14; Conservative

Matches

Local Similarity

ABK25391 standard; DNA; 17 BP.

RESULT 438

ABK25391;

ABK25391 ID ABK2 XX AC ABK2

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Male-sterile plant producing genome altering oligonucleotide #292.
                    ABK25392 standard; DNA; 17 BP
     1 creaccreaceccer 16
                                (first entry)
410 CTGAGCTGTGGGACGT
                                                                                           WO200192512-A2
                                09-APR-2002
                                                                                                 06-DEC-2001
                          ABK25392;
                                                                                     Synthetic
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Chromosomal genomic alteration; genome altering oligomucleotide; PCR; ss; o-methyl modification; LNA modification; phosphorothicate linkage; DNA repair; DNA alteration; environmental tolerance; hygromycin-B; abiotic stress tolerance; improved nutritional value; hygromycin; primer; amino acid over production; herbicide resistance; glyphosate resistance; pophyric herbicide resistance; sulphomylurea herbicide resistance; pophyric herbicide resistance; riazine resistance; disease resistance; modified oil production; modified starch production; waxy starch; modified fioral morphology; male-sterile plant; albino mutant; modified fatty acid content; reduced palmitate production; albino plant; photosynthetic process.

01-JUN-2001; 2001WO-US017672.

01-JUN-2000; 2000US-0208538P. 30-OCT-2000; 2000US-0244989P. 27-MAR-2001; 2001US-00818875.

(UYDE) UNIV DELAWARE.

Kim J; Rice MC, Gamper HB, Kmiec EB,

WPI; 2002-106307/14.

New oligonuclectides with modified nuclease-resistant termini, useful for creating plants with desired phenotypes, e.g. stress tolerance, improved nutritional value, herbicide or disease resistance, or modified oil production.

Claim 7; Page 87; 220pp; English.

The invention relates to an oligonucleotide for targeted alteration of a genetic sequence, which comprises a single-stranded oligonucleotide CC daving a DNA domain. The DNA domain has at least one mismatch with respect to the genetic sequence to be altered and further comprises chemical modifications of the oligonucleotide. The chemical modifications of the oligonucleotide. The chemical modifications of the oligonucleotide are useful for consist of o-methyl modification, an LNA modification, two or more phosphorothioate linkages on a terminus, or a combination of any two or more of these modifications. The oligonucleotides are useful for conceptides are particularly useful for content of plant genetic information. The oligonucleotides are particularly useful for creating plants with desired phenotypes, e.g. environmental or abiotic stress tolerance, improved conferring amino acid over production), herbicide resistance (e.g. altering amino acid over production), herbicide resistance (e.g. glyphosate resistance, imidazolinome and sulphonylurea herbicide.

CC glyphosate resistance, imidazolinome and sulphonylurea herbicide resistance or triazine resistance), disease resistance, modified oll production, modified starch production (e.g. increased starch or production of waxy starch), altered floral morphology (e.g. male-sterile plants) or modified fatty acid content (e.g. reduced palmitate, increased stearate or reduced linolenic acid).

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The oligonucleotides are also useful for producing albino mutants for the analysis of photosynthetic processes. This sequence represents a genome altering oligonucleotide of the invention
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                                                                                          2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 4e+02;
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                                                            Sequence 17 BP; 3 A; 7 C; 5 G; 2 T; 0 U; 0 Other;
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Matches 14; Conserv
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Human; hammerhead ribozyme; cytostatic; antitumour; antidiabetic; ophthalmological, antiarthritic; antipsoriatic; virucide; osteopathic; vulnerary; cancer; lymphoma; Ewing's sarcoma; melanoma; psoriasis; tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgaris; angiofibroma of tuberous sclerosis; port.wine stain; wound healing; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leukaemia; ss; osler-Weber-rendu syndrome, leukaemia; osteoporosis; DNAzyme; inozyme; amberzyme.

Human ERG hammerhead ribozyme target sequence, Seq ID No 839.

09-APR-2002 (first entry)

.6-MAY-2001; 2001WO-US015866. WO200188124-A2. Homo sapiens. 22-NOV-2001.

16-MAY-2000; 2000US-00572021. (RIBO-) RIBOZYME PHARM INC. (GLAX) GLAXO GROUP LTD.

Mclaughlin F, Randi AM; Jarvis T, Von Carlowitz I, Mcswiggen JA, WPI; 2002-082995/11. Novel polynucleotide which down regulates expression of Ets-related genuseful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome.

Claim 4; Page 74; 149pp; English.

The invention relates to a nucleic acid molecule (I) which down regulates expression of an Ets-related gene (ERG). (I) is useful for treating conditions selected from cancer, lymphoma, Eaving's sarcoma, melanoma, tumour angiogenesis, diabetic retinopathy, macular degeneration, neovascular glaucoma, myopic degeneration, arthritis, psoriasis, vertuca variants, simple in a syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Weber-rendu syndrome, leukaemia, osteoporosis and wound healing. (I) is useful for treatment. The method comptises the use of one or more therapies the treatment. The method comprises the use of one or more therapies under conditions suitable for the treatment. Leukaemia or tumour angiogenesis is treated by administering (I) to the patient in angiogenesis is treated by administering (I) to the patient in conjunction with one or more of other therapies such as radiation or chemotherapy treatment. (I) is useful for reducing ERG activity in a

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ERG gene, by contacting the cell with (I). (I) is useful for cleaving RNA of ERG gene, by contacting (I) with RNA, in the presence of a divalent cation such as MgA. (I) is useful for diagnosis of conditions and diseases related to the expression of ERG, and as diagnostic tool to examine genetic drift and mutations within diseased cells or to detect the presence of ERG RNA in a cell. (I) is useful for specifically targeting genes that share homology with ERG gene or ERG fusion genes. ABK17354-ABK22719 represent nucleic acids, including antisense and enzymatic nucleic acid molecules which regulate expression of ERG, and related PCR primers of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   useful for treating cancer, diabetic retinopathy, macular degeneration,
arthritis, psoriasis, verruca vulgaris and Sturge Weber syndrome.
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Matches 14; Conservative
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treating a patient having a condition associated with the level of ERG, by contacting cells of the patient with (I) under conditions suitable for the treatment. The method comprises the use of one or more therapies under conditions suitable for the treatment. Leukaemia or tumour angiogeneals is treated by administering (I) to the patient in conjunction with one or more of other therapies such as radiation or chemotherapy treatment. (I) is useful for reducing ERG activity in a cell, by contacting the cell with (I). (I) is useful for cleaving RNA of ERG gene, by contacting (I) with RNA, in the presence of a divalent cation such as Mg2+. (I) is useful for diagnosis of conditions and call make to the expression of ERG, and as diagnostic tool to examine genetic drift and mutations within diseased cells or to detect the presence of ERG RNA in a cell. (I) is useful for specifically transcribed that share homology with ERG gene or ERG fusion genes. ABK17554-ABK2279 represent nucleic acids, including antisense and contacting contacting center nucleic acids, including antisense and contacting antisense and contact
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87.5%; Pred. No. 4e+02;
tive 0; Mismatches 2; Indela
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Example 2; SEQ ID NO 716; 60pp + Sequence Listing; English

activity of human POSHL1.

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The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSH1.1) polypeptide (I), comprising a sequence of 730 amino acids (S1, ABB83999), a sequence having 65* sequence identity to (S1), (S1) having 95* deviations, especially conservative substitutions or a fragment of the sequences comprising at least 8 contiguous amino acids. Human POSH1 is a proteo-oncogene/oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as downstream components of the signal transduction pathway. (I) is useful of for identifying a specific binding partner. (I) and mucleic acids (II) encoding (I) are useful for diagnosing, monitoring disease and traating caused by altered expression of human POSH11 including diagnosing and creating cancer, they useful in the development of vaccines and (II) is useful in gene therapy. (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligomorleotide useful in examples of the invention. Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied to
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30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000667.
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30-JAN-2001; 2001WO-US000669.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABV90002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 443
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AC ABV9
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DT 23-D
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protein (POSHL 1) polypeptide (1), comprising a sequence of 730 amino acids (S1, ABBB3999), a sequence having 65% sequence identity to (S1), acids (S1, ABBB3999), a sequence having 65% sequence identity to (S1), acids (S1, ABBB3999), a sequence having 65% sequence identity to (S1), acids (S1, ABBB3999), a sequence having 65% sequence identity to (S1), acids (S1, ABBB3999), a sequence having 65% sequence identity to (S1), acids (S1, ABBB3999), a sequence of congrishing at least 8 contiguous amino acids. Human POSHL 1 is a proto-oncogene product that functions as an adaptor protein that interacts with Rho family small GTPases as well as downstream components of the signal transduction pathway. (I) is useful for identifying a specific binding partner. (I) and nucleic acids (II) for identifying a specific binding partner. (I) and nucleic acids (II) acused by altered expression of human POSHL1 including dispnosing and treating cancer, they useful in the development of vaccines and (II) is useful for constructing microarrays which are useful for measuring and for surveying gene expression and creating transgenic non-human animals capable of producing the proteins. The present sequence is that of a scanning oligonucleotide useful in examples of the invention. Note: The present sequence did not form part of the printed specification, but is based on sequence information supplied to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human leukocyte antigen (HLA) typing, useful for judging HLA genotypes of individuals e.g. by determining immunogenetic differences when transplanting between them.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a typing kit for judging human leukocyte antigen (HLA) genotype of a sample by hybridising a substrate on which 10-24 base
                                                                               The invention relates to an isolated SH3 domain (POSH)-like signalling protein 1 (POSHL 1) polypeptide (I), comprising a sequence of 730 amino APPRA 1999). a sequence having 65% sequence identity to (SI),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Moriya S, Nishida M;
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immunogenetic; transplantation; genetic disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 2.8%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 4e+02; Matches 14; Conservative 0; Mismatches 2; Indels
                                             Example 2; SEQ ID NO 715; 60pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human HLA genotyping oligonucleotide SEQ ID NO 1071.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17 BP; 3 A; 7 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Inoko H, Kagiya T, Ichihara T, Matsumura Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Derwent by the European Patent Office
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 272 CTTCTCCGGAGGCACC 287
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oligonucleotides (ABL30512-ABL31809) originating in the sequences of genes e.g. belonging to HLA class I antigens on human genome and containing gene polymorphisms as alloantigens have been immobilised as primers for amplification of cleaved nucleic acids relating to gene polymorphisms. The method is useful for judging HLA genotypes of individuals by determining immunogenetic differences before transplanting between them, providing genetic information to decide compatibility of organ and tissue for transplantation e.g. of bone marrow, kidney, liver, pancreas, Langerhans islet in pancreas and cornea, susceptibility diagnosis of genetic diseases and identifying individuals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to nucleic acid molecules that modulate replication of the West Nile Virus (WNV). The nucleic acid molecules are useful for treating a condition related to WNV infection e.g. pancreatitis, encephalitis, myocarditis, meningitis, neurologic infection, hepatitis, liver failure, hepatocellular carcinoma or cirrhosis. The nucleic acid molecule is selected from the group of ribozymes consisting of Hammerhead, Inozyme, G-cleaver, DNAzyme, Amberzyme and Zinzyme. The nucleic acid molecules further comprise at least five ribose residues, at least ten 2-0-methyl modifications, phosphorothicate linkages on at least three of the 5' terminal nucleocides and a 3' end modification of a 3' inverted abasic moiety. Nucleic acid molecules SEQ ID NO 1 to 37080
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New nucleic acid molecule that modulates replication of West Nile Virus (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                   2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 4e+02; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                         WNV minus strand Inozyme substrate SEQ ID NO 9764.
                                                                                                                                                         Sequence 17 BP; 1 A; 6 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 23; SEQ ID NO 9764; 495pp; English.
                                                                                                                                                                                                                                                                                                                                         ACN09761 standard; RNA; 17 BP
                                                                                                                                                                                                                                        379 CGGAGCGAGTCCCCGC 394
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19-OCT-2001; 2001WO-US048350.
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                                                                                                                                                                                                                                                                CGGAGCCAGTCCACGC 1
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                                                                                                                                                                                                Local Similarity 87.5
1es 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mcswiggen JA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Amberzyme; Zinzyme; ss.
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(MCSW/) MCSWIGGEN J A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-706994/76.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     West Nile Virus.
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                                                                                                                                                                                                                                                                                                                                                                     ACN09761;
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                                                                                                                                                                                    Query Match
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                                                                                                                                                                                                                                                                                                               445
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encephalitis, myocarditis, meningitis, neurologic infection, hepatitis, liver failure, hepatocellular carcinoma or cirrhosis. The nucleic acid molecule is selected from the group of ribozymes consisting of Hammerhead, Inozyme, G-leaver, DNAzyme, Amberzyme and Zinzyme. The nucleic acid molecules further comprise at least five ribose residues, at least there of the S' terminal nucleotides and a 3' end modification of least three of the S' terminal nucleotides and a 3' end modification of a 3'-3' inverted abasic moiety. Nucleic acid molecules SEQ ID NO 1 to 37080 are claimed; however, SEQ ID NO 2194-2206 and 17502-17514 are not given in the specification. The present sequence is that of a nucleic acid
are claimed; however, SEQ ID NO 2194-2206 and 17502-17514 are not given in the specification. The present sequence is that of a nucleic acid molecule of the invention
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                                                                                                                                                                                                                      DB 1; Length 17;
                                                                                                                                                                                                                                                                                           Indels
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                                                                                                                                                 G; 0 T; 5 U; 0 Other,
                                                                                                                                                                                                               2.8%; Score 12.8; DB 1
52.5%; Pred. No. 4e+02;
ve 4; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WNV Zinzyme substrate SEQ ID NO 4595.
                                                                                                                                                                                                                                                                                                                                                            316 TCAGCCGCGGGTCTCT 331
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP
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1 UGAGCCGCAGGUCUCU 16
                                                                                                                                                 Sequence 17 BP; 2 A; 5 C; 5
                                                                                                                                                                                                                                                          62.5%;
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                                                                                                                                                                                                                                                   Local Similarity 62.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-706994/76.
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                                                                                                                                                                                                                                                                                               Matches
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wnv; west Nile Virus; antiinflammatory; cytostatic; hepatotropic; virucide; neuroprotective; antibacterial; replication; pancreatitis; encephalitis; myocarditis; meningitis; infection; hepatitis; litter failure; cancer; Cirrhosis; Hammerhead; Inozyme; DNAzyme;

Amberzyme; Zinzyme; ss

West Nile Virus. WO200268637-A2. 06-SEP-2002

20-OCT-2000; 2000US-0242411P. 19-OCT-2001; 2001WO-US048350.

RIBO-) RIBOZYME PHARM INC

(MCSW/) MCSWIGGEN J A.

BLAT/)

Mcswiggen JA;

Blatt L,

WNV Hammerhead Ribozyme substrate SEQ ID NO 405.

22-APR-2004 (first entry)

ACN00415;

ACN00415 standard; RNA; 17 BP.

RESULT 448

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ACN00415,

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of the West Nile Virus (WNV). The nucleic acid molecules are useful for treating a condition related to WNV infection e.g. pancreatics, encephalitis, meningitis, neurologic infection, hepatitis, inversional infection e.g. pancreatics, liver failure, hepatocellular carcinoma or cirrhosis. The nucleic acid molecule is selected from the group of ribozymes consisting of hammerhead, inozyme, G-cleaver, DNAzyme, Amberzyme and Zinzyme. The nucleic acid molecules further comprise at least five ribose residues, at least ten 2'-0-methyl modifications, phosphorothicate linkages on at least three of the 5' terminal nuclectides and a 3' end modification of a 3'-3' inverted abasic moiety. Nucleic acid molecules SEQ ID NO 1 to 37080 are claimed; however, SEQ ID NO 2194-2206 and 17502-17514 are not given in the specification. The present sequence is that of a nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to nucleic acid molecules that modulate replication
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New nucleic acid molecule that modulates replication of West Nile Virus (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
                       Gaps
                                                                                                                                                                                                                                                                                                                                     WNV, West Nile Virus, antiinflammatory; cytostatic; hepatotropic; viruoids; neuroprotective; antibacterial; repplication; panoreatitis; encephalitis; myocarditis; meningitis; infection; hepatitis; liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;
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                     Indels
                                                                                                                                                                                                                                                                                                    WNV minus strand Amberzyme substrate SEQ ID NO 15002.
                   2;
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87.5%; Pred. No. 4e+02;
ive 0; Mismatches
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                                                                                                                                                                                     ACN14999 standard; RNA; 17 BP.
                                                      316 TCAGCCGCGGGTCTCT 331
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                                                                                         16 rgagccgcaggrerer 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       molecule of the invention
                   14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Blatt L, Mcswiggen JA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (BLAT/) BLATT L.
(MCSW/) MCSWIGGEN J A.
                                                                                                                                                                                                                                                                                                                                                                                                                   Amberzyme; Zinzyme; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-706994/76.
 Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                  West Nile Virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200268637-A2.
                                                                                                                                                                                                                                                               22-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              36-SEP-2002
                                                                                                                                                                                                                          ACN14999;
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                                                                                                                                                    447
                   Matches
                                                                                                                                                                     ACN14999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New nucleic acid molecule that modulates replication of West Nile Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17 BP; 6 A; 2 C; 7 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 23; SEQ ID NO 405; 495pp; English.
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ID ACN03272 standard, RNA; 17 |
XX AC ACN03272;
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Best Local Similarity 87.5
Matches 14; Conservative
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Gaps

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TCTCGCTGACTTTCAG 118

103

Conservative

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liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;
                                    West Nile Virus
                                                            #0200268637-A2
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                                                                                    06-SEP-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                        of the West Nile virus (WNV). The nucleic acid molecules are useful for treating a condition related to WNV infection e.g. pancreatitis, encephalitis, myocarditis, meningitis, neurologic infection, hepatitis, liver failure, hepatocellular carcinoma or cirrhosis. The nucleic acid molecule is selected from the group of ribozymes consisting of hammerhead, Inozyme, G-Cleaver, DNAzyme, Amberzyme and Zinzyme. The nucleic acid molecules further comprise at least five ribose residues, at least three of the 5' terminal nuclectides and a 3' end modification of a 3'-3' inverted abasic moiety. Nucleic acid molecules SEQ ID NO 194-2206 and 17502-17514 are not given in the specification. The present sequence is that of a nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to nucleic acid molecules that modulate replication
                                                                                                                                                                                                                                                                                                                                                     New nucleic acid molecule that modulates replication of West Nile Virus (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                               WNV, West Nile Virus, antiinflammatory; cytostatic; hepatotropic; virucide; neuroprotective; antibacterial; replication; pancreatitis; encephalitis; myocarditis; meningitis; infection; hepatitis; liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WNV; West Nile Virus; antiinflammatory; cytostatic; hepatotropic; virucide; neuroprotective; antibacterial; replication; pancreatitis; encephalitis; myocarditis; meningitis; infection; hepatitis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 2.8%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 4e+02; Matches 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WNV minus strand DNAzyme substrate SEQ ID NO 14013.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 5 A; 5 C; 5 G; 0 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                      Claim 23; SEQ ID NO 3275; 495pp; English.
                        WNV Inozyme substrate SEQ ID NO 3275.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                316 TCAGCCGCGGGTCTCT 331
                                                                                                                                                                                                  19-OCT-2001; 2001WO-US048350.
                                                                                                                                                                                                                          20-OCT-2000; 2000US-0242411P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17 TGAGCCGCAGGTCTCT 2
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                                                                                                                                                                                                                                                    RIBOZYME PHARM INC.
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(first entry)
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                                                                                                                                                                                                                                                                                                    Blatt L, Mcswiggen JA;
                                                                                                   Amberzyme; Zinzyme; ss
                                                                                                                                                                                                                                                                            (MCSW/) MCSWIGGEN J A.
                                                                                                                                                                                                                                                                                                                            WPI; 2002-706994/76.
                                                                                                                        West Nile Virus.
                                                                                                                                                 WO200268637-A2
 22-APR-2004
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of the West Nile Virus (WNV). The nucleic acid molecules are useful for treating a condition related to WNV infection e.g. pancreatitis, encephalitis, myocarditis, meningitis, neurologic infection, hepatitis, meningitis, neurologic infection, hepatitis, molecule is selected from the group of ribozymes consisting of Hammerhead, Inozyme, G-cleaver, DNAzyme, Amberzyme and Zinzyme. The nucleic acid molecules further comprise at least five ribose residues, at least three of the 5' terminal nucleocides and a 3' end modification of a 3'-3' inverted abasic molety. Nucleic acid molecules SEQ ID No 194-2206 and 17502-17514 are not given in the specification. The present sequence is that of a nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to nucleic acid molecules that modulate replication
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New nucleic acid molecule that modulates replication of West Nile Virus (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Seguence 17 BP; 2 A; 6 C; 3 G; 0 T; 6 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human MDZ3 scanning oligonucleotide SEQ ID 815.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 23; SEQ ID NO 14013; 495pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           103 TCTCGCTGACTTTCAG 118
                                                                                                                                                                                                                                                                                               19-OCT-2001; 2001WO-US048350.
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1 UCUCUCUGACCUUCAG 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                        (RIBO-) RIBOZYME PHARM INC. (BLAT/) BLATT L. (MCSW/) MCSWIGGEN J A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADA99826 standard; DNA; 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Blatt L, Mcswiggen JA;
Amberzyme; Zinzyme; ss
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Best Loca
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is cancoded at chromosome 7422.1, MDZ4 is encoded at chromosome 6721.3-22.2, MDZ7 is encoded at chromosome 6721.3-22.2, MDZ7 is encoded at chromosome 1601.2 and MDZ12 is encoded at chromosome 572.1, mDZ7 is encoded at chromosome 1601.2 and MDZ12 is encoded at chromosome 582.2, mDZ7, MDZ7, MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are useful as therapeutic agents for gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
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                                                                                                                                                                                                                                                                                                            New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ7 or MDZ12, e.g. cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human MDZ3 scanning oligonucleotide SEQ ID 813.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 8; SEQ ID NO 815; 103pp; English.
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                             30-JUL-2002; 2002EP-00016874.
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Best Local Similarity 87.5%;
Marches 14; Conservative
                                                                                02-AUG-2001; 2001US-00922181
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                                                                                                                                                                                                 Gu Y, Nguyen C;
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                                                                                                                                           (AEOM-) AEOMICA INC
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                                                                                                                                                                                                 Shannon M,
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The present invention relates to novel human zinc finger-containing proteins and their coding sequences: MDZ3, MDZ4, MDZ12, MDZ12. MDZ3 is cancoded at chromosome 7422.1, MDZ4 is encoded at chromosome 6p21.3-22.2, MDZ7 is encoded at chromosome 16p21.2 and MDZ12 is encoded at chromosome 16p2.3 and MDZ12 is encoded at chromosome 16p2.3 and MDZ12 is encoded at chromosome 15q26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 are useful in therapy, or MDZ3, MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for diagnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agente for gene therapy or as proteins are useful as therapeutic agente for gene therapy or as
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                                               New zinc finger-containing proteins and nucleic acids, useful in manifacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ4, MDZ7 or MDZ12, e.g. cancer.
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                                                                                                                                                                                      Example 8; SEQ ID NO 813; 103pp; English.
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WPI; 2003-423107/40.
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ABZ61267;
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                                                               acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule invention has cytostatic, anti-HIV, and anti-theumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABESS989 - ABES62216, ABZ6444 - ABZ65531, ABZ66520 - ABZ66524, ABZ66530 - ABZ66530 - ABZ66520 - ABZ66524, ribozymes of the invention
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                                                The invention relates to a novel short interfering RNA (siRNA) nucleic
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enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; ss.
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  Claim 58; Page 129; 185pp; English
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06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
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Matches 14; Conservative
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shown in ABZ59889 - ABZ65216, ABZ64544 - ABZ65531, ABZ66520 - ABZ66524,
ABZ66530 - ABZ66585 represent substrate/target sequences for the human
ribozymes of the invention
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                                                                                                                                 Seguence 17 BP; 2 A; 11 C; 1 G; 0 T; 3 U; 0 Other;
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06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
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Best Local Similarity 87.5%
Local 14; Conservative
                                                                                                                                                                                                                                                                                                                  6 GCGGAGGGTGGGCCTG
                                                                                                                                                                                                                                                                                                                                                                              17 GAGGAGGGTGGGGCTG
                                                                                                                                                                                                                      Local Similarity 87.9
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Human H-Ras DNAzyme target #179.
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                                                                                                                                                                                                                                                                                                                                                                                                                 Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HBR2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
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                                                                                                                                                           Human, ribozyme, short interfering RNA, siRNA; HER2, K-Ras;
enzymatic nucleic acid, H-Ras; N-Ras, HIV, cytostatic, anti-HIV;
anti-rheumatic; cancer, AIDS; ss.
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87.5%; Pred. No. 4e+02;
Live 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 4; Page 133; 185pp; English.
                                                                                                                                      Human HER2 DNAzyme substrate #20.
                                                                   ABZ64563 standard; RNA; 17 BP
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          16 GCCGCGGTCGGGCGGG 1
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                                                                                                                 (first entry)
Local Similarity 87.5 les 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-140484/13.
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                                                                                                                                                                                                           Homo sapiens.
                                                                                                                 21-MAR-2003
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The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HER2, K.Ras, H.Ras, N.Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-HER2, K.Ras, H.Ras, and HIV activity in a cell. The nucleic acid stream the same useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABES9899 - ABES62216, ABE65531, ABE66520 - ABE66524, ribozymes of the invention
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HBV reverse transcriptase, Enhancer I region, viral replication,
degenerative, disease state, HBV infection, HCV infection, cirrhosis,
liver failure, hepatocellular carcinoma, hepatotropic, cytostatic,
virucide, antiinflammatory, substrate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HBR2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
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Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras;
enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV;
anti-rheumatic; cancer; AIDS; ss.
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10-SEP-2001; 2001US-0318471P.
                                                                                                                                                                                                                                                                                                                                                   29-MAY-2001; 2001US-0294140P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            202 TCCCGGGGACCTGCGG 217
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                                                                                                                                                                                                                                                                                            29-MAY-2002; 2002WO-US016840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 UCCCUGGGGCCUGCGG 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 75.0
nes 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-140484/13.
                                                                                                                                                                            WO200297114-A2
                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                   05-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mcswiggen J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-SEP-2003
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21 GGGAGGGTGGTGGCC 36

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ABZ61388 standard; RNA; 17

RESULT 457

ABZ61388

ABZ61388;

HXXXX

21-MAR-2003 (first entry)

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Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
                                     Nucleic acid molecule, Hepatitis C virus, HCV; Hepatitis B virus, HBV; RNA stability; RNA expression; RNA synthesis, antisense; enzymatic nucleic acid; hammerhead ribozyme; DNAzyme; inozyme; zinzyme;
                                                                          amberzyme; G-cleaver ribozyme; decoy molecule; apramer;
HBV reverse transcriptase; Enhancer I region; viral replication;
degenerative; disease state; HBV infection; HCV infection; cirrhosis;
liver failure; hepatocellular cardinoma; hepatotropic; cytostatic;
            minus strand DNAzyme substrate sequence #1478.
                                                                                                                                  virucide; antiinflammatory; substrate; ss.
                                                                                                                                                                                                                                                                     26-MAR-2001; 2001US-00817879.
08-JUN-2001; 2001US-00877478.
08-JUN-2001; 2001US-0296876P.
24-OCT-2001; 2001US-0337055P.
05-DEC-2001; 2001US-0337055P.
                                                                                                                                                                                                                                             26-MAR-2002; 2002WO-US009187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-229207/22
                                                                                                                                                             Hepatitis C virus
                                                                                                                                                                                         WO200281494-A1
                                                                                                                                                                                                                  17-0CT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            infection.
                                                                                                                                                                                                                                                                                                                                                                                                                           (PAVC/)
(LEEP/)
(DRAP/)
(ROBE/)
                                                                                                                                                                                                                                                                                                                                                        (RIBO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Draper
                                                                                                                                                                                                                                                                                                                                                                                                              (MORR/)
                                                                                                                                                                                                                                                                                                                                                                      (BLAT/
                                                                                                                                                                                                                                                                                                                                                                                                 MCSW/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Blatt
The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or and enzymatic nucleic acids such as harmerhead molecules include antisense and enzymatic nucleic acids such as harmerhead ribozymes, DNAzymes, inozymes, zinzymes, amberzymes, and G-cleaver ribozymes, Also disclosed transcriptase and/or HBV reverse transcriptase primer sequences, as well as Oligonucleotides that specifically bind the Enhancer I region of HBV genes and Alox people and Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression of HBV compounds and/or potential therapies directed against HBV, and compounds compounds and/or potential therapies directed against HBV, and compounds methods of the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene carcinoma. The present sequence represents a substrate for one of the HCV interest of the compounds and carcinoma. The present sequence represents a substrate for one of the HCV interest.
                                                                                                                                                                                                                                                                                                                                                                                                              Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
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                                                                                                                                                                                                                                                                                                                                             Lee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ? Match 2.8%; Score 12.8; DB 1; Length 17; Local Similarity 75.0%; Pred. No. 4e+02; les 12; Conservative 2; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                           Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17 BP; 1 A; 1 C; 11 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                           Morrissey D,
                                                                                                                                                                                                                                                                                                                                           Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 259; 387pp; English.
                                                                                                                                  2001US-00877478.
2001US-0296876P.
                                                                                                                                                            24-OCT-2001; 2001US-0335059P.
                                                                                           26-MAR-2002; 2002WO-US009187
                                                                                                                                                                                                     RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                          Macejak D,
Roberts E;
                                                                                                                                                                                                                               MACEJAK D.
MCSWIGGEN J.
MORRISSEY D.
                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-229207/22.
                                                                                                                                                                                                                                                                                                                ROBERTS E.
           Hepatitis C virus
                                                                                                                                                                                                                                                                                     LEE P.
DRAPER K.
                                                                                                                                                                                                                                                                          PAVCO P.
                                      WO200281494-A1
                                                                                                                                                                                                                    BLATT
                                                                                                                                  08-JUN-2001;
                                                                                                                      26-MAR-2001;
                                                                 17-OCT-2002
                                                                                                                                                                                                                                                                                                                                          Blatt L, M
Draper K,
                                                                                                                                                                                                                                                                                                                                                                                                                                         infection.
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                                                                                                                                                                                                                                                         (MORR/) 1
(PAVC/) 1
(LEEP/) 1
(DRAP/) 1
(ROBE/) 1
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                                                                                                                                                                                                    (RIBO-)
(BLAT/)
(MACE/)
(MCSW/)
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RIBOZYME PHARM INC

MACEJAK D. MCSWIGGEN J. MORRISSEY D.

ä Lee

ď Pavco

Mcswiggen J, Morrissey D,

Macejak D, Roberts E;

о п, , к,

ROBERTS E DRAPER K. PAVCO P.

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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, amberzymes, amberzymes, and G-cleaver ribozymes, Also disclosed re nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptase and/or HBV reverse transcriptase primer sequences, as well as oligomucleotides that specifically bind the Enhancer I region of HBV genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or lapication and/or percential therapies directed against HBV, and compounds methods of the invention are useful for the treatment of degenerative and elsease states related to HBV and HCV infection, replication and gene carcinoma. The present sequence represente a substrate for one of the HCV DNAzyme or minus strand DNAzyme sequences disclosed in the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 4e+02; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 17 BP; 1 A; 6 C; 6 G; 0 T; 4 U; 0 Other;
Claim 1; Page 301; 387pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 07.59
Marches 14; Conservative
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Gaps

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32

20 TGGGAGGGGTGGTGGC 12; Conservative

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Matches

2 UGGGAGGGUGGGUGGC 17

ACD64287 standard; RNA; 17

RESULT 459

(first entry)

30-SEP-2003

ACD64287;

ACD64287/c ID ACD64 XX AC ACD64 XX DT 30-SE

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Gaps

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433 GGACTCGGCTCACACA 448

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the synthesis, expression and/or stability of Hepatitis C virus (HCV) or
Hepatitis B virus (HHV) RNA. The nucleic acid molecules include antisense
and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes,
inozymes, zinzymes, amberzymes, and G-cleaver ribozymes. Also disclosed
are nucleic acid decoy molecules and aptamers that bind to HBV reverse
transcriptase and/or HBV reverse transcriptase primer sequences, as well
as oligonucleotides that specifically bind the Enhancer I region of HBV
DNA. The nucleic acids may be used to modulate the expression of HBV
genes and HBV viral replication. Also disclosed is a method for screening
compounds and/or potential theraphse directed against HBV, and compounds
that modulate the expression and/or replication of HCV. The compounds and
methods of the invention are useful for the treatment of degenerative and
disease states related to HBV and HCV infection, replication and gene
expression such as cirrhosis, liver failure, and hepatocellular
                                                                                                                                                                                                                                                                                                                                                                                                  Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; hammerhead ribozyme; DNAzyme; inozyme; zinzyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Lee P;
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                                                                                                                                                                                                                                                                                                                                             HCV minus strand DNAzyme substrate sequence #1362.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 1; Page 299; 387pp; English.
                                                                                                                                                                               BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-MAR-2001; 2001US-00817879.
08-JUN-2001; 2001US-00877478.
08-JUN-2001; 2001US-0296876P.
24-OCT-2001; 2001US-0337055P.
05-DEC-2001; 2001US-0337055P.
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                                                                                                                                                                         ACD64059 standard; RNA; 17
(first entry)
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Roberts E;
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MCSWIGGEN J.
MORRISSEY D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-229207/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hepatitis C virus.
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DRAPER K.
ROBERTS E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PAVCO P.
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                                                                                                                                                                                                                                  ACD64059;
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(BLAT/)
(MACE/)
(MCSW/)
(MORR/)
(PAVC/)
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(ROBE/)
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carcinoma. The present sequence represents a substrate for one of the HCV
DNAzyme or minus strand DNAzyme sequences disclosed in the present
invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                          Nucleic acid molecule, Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid, hammerhead ribozyme; DNAzyme; inozyme; zinzyme; amberzyme; G-claever ribozyme; decoy molecule; aptemer; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
                                                                                                                                     Gaps
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                                                                                                 2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 4e+02; ative 0; Mismatches 2; Indels
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                                                                       Sequence 17 BP; 3 A; 6 C; 7 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Morrissey D,
                                                                                                                                                                                                                                                                                                                                                  HBV inozyme substrate sequence #352.
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                                                                                                                                                            208 GGACCTGCGGCGGGTC 223
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08-JUN-2001; 2001US-00877478.
08-JUN-2001; 2001US-0296876P.
24-CCT-2001; 2001US-0335059P.
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                                                                                                                                                                                                                                                             ACD52324 standard; RNA; 17
                                                                                                                                                                              GCACCTGCGGCGGCTC
                                                                                                                                                                                                                                                                                                                      (first entry)
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                                                                                                                                Conservative
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Roberts E;
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MACEJAK D.
MCSWIGGEN J.
MORRISSEY D.
                                                                                                                 Local Similarity
les 14; Conserv
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LEE P.
DRAPER K.
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Draper K,
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                                                                                                                                                                                                                                                                                           ACD52324;
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                                                                                                 Query Match
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(MACE/)
(MCSW/)
(MORR/)
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(LEEP/)
(DRAP/)
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                                                                                                                  Best Loca
Matches
                                                                                                                                                                                                                                RESULT 461
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inozymes, zinzymes, amberzymes, and G-cleaver ribozymes. Also disclosed are nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptase and/or HBV reverse transcriptase primer sequences, as well as oligonuclectides that specifically bind the Enhancer I region of HBV genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HV. The compounds of the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene expression such as cirrhosis, liver failure, and hepatocellular carcinoma. The present sequence represents a substrate for one of the HBV ribozyme, inozyme, G-cleaver, zinzyme, DNAzyme or amberzyme sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to human phenotype-associated (PA) genes (I ; Appl3307-AbF13386) which contain a Single Nuclectide Polymorphism (SNP). The SNP is given in the sequence as a variant nucleotide Also claimed are methods for screening for agents which regulate the activity of a PA gene and reagents that modulate the activity of a PA purcleotide where the reagent is identified by the screening methods. The methods and compositions of the present invention are useful for prognosticating, diagnosing and treating cardiovascular diseases. Mypertension, restenosis, arterial inflammation, mayocardial infarction and stroke. The present sequence is a PCR primer, used in the examples from the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cardiant; antiarteriosclerotic; vasotropic; cerebroprotective;
hypotensive; gene therapy; human; SNX9; Sorting Nexin 9; PCR; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                   Gape
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New isolated polynucleotide encoded by a phenotype-associated gene, useful for prognosticating statin therapy response, and diagnosing treating cardiovascular diseases, such as hypertension, myocardial
                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                        2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 4e+02; ive 0; Mismatches 2; Indel8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SNX9 (Sorting Nexin 9), BaySNP 6743, PCR primer #2.
                                                                                                                                                                                                                                                                                                   Sequence 17 BP; 1 A; 8 C; 4 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Page 69; 182pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               468/c
ADF13468 standard; DNA; 17 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                      117 AGCGGGGGAAAAGCC 132
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               16 AGCGGGCGGTAGAGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                               14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Schwers S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   infarction and stroke.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-712738/67.
                                                                                                                                                                                                                                                                                                                                                             Similarity
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                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 462
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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5, or 3, end genomic flanking regions, 5 and 3 intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an entitinflammatory servoid and ubiquinone. A composition of the invention has antiinflammatory servoid and ubiquinone, A composition of the invention has antiinflammatory, antiallargic, antiasthmatic, hypotensive, communosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of or reducing benefitivity to adenosine, reducing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at fip. Wipo: Allery wipo at fip. WIPO at the sequences.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human, antisense, lung dysfunction, nasal airway dysfunction, antiinflammatory steroid, ubiquinone, antiinflammatory; antiallergic; antiathamatic; hypotensive; fromtunesuppressive, cytostartic; gene therapy; antisense gene therapy; respiratory, lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                    Gaps
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                                                      Query Match 2.8%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 4e+02; Matches 14; Conservative 0; Mismatches 2; Indel8
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                 Sequence 17 BP; 1 A; 11 C; 2 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; SEQ ID NO 11632; 872pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   lung inflammation; respiratory disease; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                     Human C/EBP antisense fragment no.2250.
                                                                                                                                             334 GGGGCGAGGCGAGGT 349
                                                                                                                                                                                                                                                                                           ВР
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                                           ABZ96390 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                       GGGCGAGGCCAGGGT
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Tang L,
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Miller S,
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                                                                                                                                                                                                                                                    RESULT 463
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Sequence 17 BP; 2 A; 5 C; 8 G; 0 T; 2 U; 0 Other;

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The invention comprises nucleic acids (e.g. antisense oligonucleotides)
that down regulate the expression or inhibit the function of a receptor
for a neurite growth inhibitor, NOGO, prostaglandin D2 receptor (PTGDR),
KappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the
invention are useful for treating; cerebrovascular accident, central
nervous system (CKS) injury, spinal cord injury, cancer (e.g. melanoma,
lymphoma or glioma), inflammatory disease (e.g. rhematoid arthritis,
restenosis or asthma), Crohn's disease, diabetes, obesity, autoimmune
disease, lupus, multiple sclerosis, transplant/graft rejection,
ischemeia/reperfusion injury, glomerulonephritis, sepsis, and allergic
conditions (e.g. asthma, allergic rhinitis or atopic dermatitis). The
nucleic acids of the invention are also useful for down-regulating the
expression of a target gene and as a diagnostic tool to examine genetic
drifts and mutations within diseased cells or to detect the presence of a
target RNA in a cell. The present RNA sequence represents a human IKK-
gamma substrate sequence.
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                                                                      ch 2.8%; Score 12.8; DB 1; Length 17;
1 Similarity 87.5%; Pred. No. 4e+02;
14; Conservative 0; Mismatches 2; Indels
                                                                                                                                                    Indels
Sequence 17 BP; 0 A; 12 C; 5 G; 0 T; 0 U; 0 Other;
                                                                                                                                                    2
                                                                                                                                            0; Mismatches
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                                                                                                                                                                                                                      257 GCGGTCGGCCCGGGGC 272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADL48687 standard; RNA; 17 BP.
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29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
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                                                                                                             Local Similarity
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                                                                      Query Match
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                                                                                                                                        Matches
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The invention comprises nucleic acids (e.g. antisense oligonucleotides)
that down regulate the expression or inhibit the function of a receptor
for a neurite growth inhibitor, NOGO, prostaglandin Dz receptor (PTGDR),
KappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the
invention are useful for treating: cerebrovascular accident, central
nervous system (CNS) injury, spinal cord injury, cancer (e.g. malanoma,
lymphoma or glioma), inflammatory disease (e.g. rheumatoid arthritis,
restenosis or asthma), Crohn's disease, diabetes, obesity, autoimmune
ischaemia/reperfusion injury, glomerulonephritis, sepsis, and allergic
conditions (e.g. asthma, allergic rhinitis or atopic dermatitis). The
nucleic acids of the invention are also useful for down-regulating the
expression of a target gene and as a diagnostic tool to examine genetic
drifts and mutations within diseased cells or to detect the presence of a
character of the present RNA sequence represents a human PKR
                                                                                                                                                                                                                                                                                                                                                                          antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restemoisis, asthma; Crohn; adiasase; diabetes; obesity; autoimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepais; allergic allergic rhinitis; atopic dermatitis; human PTGDR;
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                                               Gaps
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  2.8%; Score 12.8; DB 1; Length 17; 75.0%; Pred. No. 4e+02; 1ive 2; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                         Human PTGDR substrate sequence #681.
                                                                                 246 GCCTGGAGGCCGCGT 261
                                                                                                                                                                                                                    ADL51562 standard; RNA; 17 BP.
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29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
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                                                                                                                                                                                                                                                                                                   (first entry)
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                                         12; Conservative
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Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              substrate; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unidentified.
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                                                                                                                                                                                                                                                            ADL51562;
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                                         Matches
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                                                                                                                                                 RESULT 467
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                                                                                                                                                                                                                                                                                     antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restenosis; asthma; Crohn's disease; diabetes; obesity; autorimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis; allergy; asthma; allergic rhinitis; atopic dermatitis; human PTGDR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.
                                                      Gaps
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                           DB 1; Length 17;
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                                                      2; Indels
Sequence 17 BP; 0 A; 6 C; 8 G; 0 T; 3 U; 0 Other;
                         2.8%; Score 12.8; DB 18.8%; Pred. No. 4e+02; ve 3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 161; SEQ ID NO 5060; 317pp; English
                                                                                                                                                                                                                                                              Human PTGDR substrate sequence #646.
                                                                                182 GCTGCTGGCCCGTTCG 197
                                                                                                                                                                             ADL51527 standard; RNA; 17 BP
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29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
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                                        68.88;
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                                                                                                                                                                                                                                    (first entry)
                                        Best Local Similarity 68.8
Matches 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Blatt L, Chowrira B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-058513/05.
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                                                                                                                                                                                                                                                                                                                                                                                                                   substrate; ds
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                                                                                                                                                                                                                                   20-MAY-2004
                                                                                                                                                                                                         ADL51527;
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                           Query Match
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The invention comprises nucleic acids (e.g. antisense oligonucleotides)
that down regulate the expression or inhibit the function of a receptor
for a neurite growth inhibitor, NOGO, prostaglandin DZ receptor (PTGDR),
is papal kinase (IKK), or protein kinase PKR. The nucleic acids of the
invention are useful for treating: cerebrovascular accident, central
nervous system (CNS) injury, spinal cord injury, cancer (e.g. melanoma,
lymphoma or glioma), inflammatory disease (e.g. rheumatoid arthritis,
restenosis or aethma), Crohn's disease, diabetes, obesity, autoimmune
disease, lupus, multiple sclerosis, transplant/graft rejection,
is chaemia/reperfusion injury, glomerulonephritis, sepsis, and allergic
conditions (e.g. asthma, allergic rhinitis or atopic dermatitis). The
nucleic acids of the invention are also useful for down-regulating the
expression of a target gene and as a diagnostic tool to examine generate
carget RNA in a cell. The present RNA sequence represents a human NOGO
receptor amberzyme substrate sequence.
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rheumatoid arthritis;
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                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          graft rejection; ischaemia; reperfusion; glomerulonephritis; sepsis; allergy; asthma; allergic rhinitis; atopic dermatitis;
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                                                                         2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 4e+02; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human NOGO receptor amberzyme substrate sequence #59.
G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                    389 CCCCGCGCGCGCGC 404
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29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
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    Sequence 17 BP; 2 A; 7 C; 8
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                                                    Query Match
Best Local Similarity 87.5'
                                                                                                                                                                                                                                                                                                                 CTCCGCGCGCGGCTCG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-058513/05.
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The invention comprises nucleic acids (e.g. antisense oligonucleotides)
that down regulate the expression or inhibit the function of a receptor
for a neurite growth inhibitor, NOGO, prostaglandin Dz receptor (FOTOR),
IkappaB kinase (IKK), or protein kinase PKR. The nucleic acids of the
invention are useful for treating: cerebrovascular accident, central
nervous system (CNS) injury, spinal cord injury, cancer (e.g. melanoma,
C lymphoma or glioma), inflammatory disease (e.g. rheumatoid arthritis,
restenosis or asthma), Crohn's disease, diabetes, obesity, autoimmune
disease, lupus, multiple sclerosis, transplant/graft rejection,
ischemia/repetfusion injury, glomerulonephritis, sepsis, and allergic
conditions (e.g. asthma, allergic rhinitis or atopic dermatitis). The
nucleic acids of the invention are also useful for down-regulating the
expression of a target gene and as a diagnostic tool to examine genetic
driftes and mutations within diseased cells or to detect the presence of a
target RNA in a cell. The present RNA sequence represents a human IKK-
gamma substrate sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                         antisense oligonucleotide; neurite growth inhibitor; NOGO; prostaglandin D2 receptor; PTGDR; IkappaB kinase; IKK; protein kinase PKR; cerebrovascular accident; central nervous system injury; CNS injury; spinal cord injury; cancer; melanoma; lymphoma; glioma; inflammatory disease; rheumatoid arthritis; restenosis; asthma; Crohn's disease; diabetes; obesity; autoimmune disease; lupus; multiple sclerosis; transplant rejection; graft rejection; ischaemia; reperfusion; glomerulonephritis; sepais; allergy; asthma; allergic rhinitis; atopic dermatitis; Human IKK-gamma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel enzymatic nucleic acid that down-regulates expression of neurite growth inhibitor receptor, prostaglandin D2 receptor, IkappaB kinase or protein kinase PKR genes, for treating cancer and inflammatory disease.
                                                                                  Gaps
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                                          DB 1; Length 17;
                                                                                  2; Indels
Sequence 17 BP; 0 A; 6 C; 8 G; 0 T; 3 U; 0 Other;
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                                      2.8%; Score 12.8; DB 1
75.0%; Pred. No. 4e+02;
tive 2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                       Human IKK-gamma substrate sequence #376.
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                                                                                                                         263 GGCCCGGGGCTTCTCC 278
                                                                                                                                                                                                                                                                   ADL47866 standard; RNA; 17 BP
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29-MAY-2001; 2001US-0294412P.
28-AUG-2001; 2001US-0315315P.
                                                                                                                                                  2 GGCCCGGGCCUGUUCC 17
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                                                                                12; Conservative
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                                                           Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nucleic acid molecule in a manner that allows its expression), a mammalian cell including the expression vector and an enzymatic nucleic acid molecule that cleaves RNA derived from a GRID gene. The nucleic acid molecule is useful for treating a condition associated with the level of GRID, e.g. tissue/graft rejection and leukaemia. The present sequence is a target region for the enzymatic nucleic acids of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New nucleic acid molecule that down-regulates expression of Grb2-related with insert domain (GRID) gene, useful for treating a condition associated with the level of GRID, e.g. tissue/graft rejection and
                                                                                                                                                                                                                                                                                                                                                                                                                                   NCH ribozyme; G-cleaver ribozyme; Zinzyme; DNAzyme; amberzyme; Inozyme; hairpin ribozyme; tissue rejection; graft rejection; leukaemia.
                                                                                                                                                                                                                                                                                                                                                                                                                   Human; ss; GRID; Grb2-related with insert domain; hammerhead ribozyme;
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                                                     DB 1; Length 17;
                                                                                            Indels
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                                                                                          2;
                 Sequence 17 BP; 1 A; 6 C; 8 G; 0 T; 2 U; 0 Other;
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                                                                      4e+02;
                                                   2.8%; Score 12.8; Dilarity 75.0%; Pred. No. 4e+0 Conservative 2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                            Human GRID mRNA substrate sequence #331.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 4; SEQ ID NO 331; 74pp; English.
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                                                                                                                                                                                                                                                               ADM54056 standard; mRNA; 17
                                                                                                                             249 TGGAGGCCGCGGTCGG
                                                                                                                                                                                                                                                                                                                                        (first entry)
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MCSWIGGEN J.
HAMBLIN P A.
                                 Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               JARV/) JARVIS T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US2003134806-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                        03-JUN-2004
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                                                                                                                                                                                                                                                                                                     ADM54056;
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(MCSW/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antialergic; antiinflammatory; antialantacic; analysesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ds.
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                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antisense; bronchoconstriction; allergy; hyposecretion; pain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted toucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pabalan J, Aguilar D;
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2.8%; Score 12.8; DB 1; Length 17;
87.5%; Pred. No. 4e+02;
ative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 15; SEQ ID NO 11632; 763pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    s,
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Tang L, Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                                                                   Human C/EBPN DNA fragment 2250.
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                                                                                                 CCTCCCGGGGACCTGC 215
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                                                                                                                                                                                                                                                  1299/c
ABD20299 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                Conservative
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                      Similarity
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                                                                                                                                                                                                                                                                                                                       ABD20299;
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Query Match
Best Local
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inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gape
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            marker; GEM;
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                                                                                                                                                                                                                                                                                                                                                                                                                              2.8%; Score 12.8; DB 1; Length 17;
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0; Mismatches
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01-APR-2003; 2003US-0458978P.
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nes 14; Conserv
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Disclosure, SEQ ID NO 1053; 122pp; English.

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dentifying test compound as a potential anticancer drug if it decreases the expression of at least one gene; (3) identifying (M4) at test compound as a potential anticancer or antiglioma drug involves contacting a test compound with the cell which expresses mRNA of at least one gene; compound with the cell which expresses mRNA of at least one gene identifying the test compound as a potential anticancer drug if it decreases the expression of at least one gene; and identifying the test compound as a potential anticancer drug if it decreases the expression of at least one gene; and (4) inducing (M5) an immune response to glioma involves administering to a mammal, a protein or (1). (1) have cytostatic activities, and can be used to trigger immune destruction of glioma cells, and as immune response inducers. (M1) is useful for aiding in diagnosing glioma. (M2) is useful for treating multicature glioma in a human. (M3) is useful for inducing an immune response to a glioma in a human. (M3) is useful for inducing an immune response to a glioma in a mammal having glioma or in a mammal who has had a glioma surgically removed. The present sequence represents a human GEM content and a glioma cells. (M2) is useful for inducing an immune response to a glioma in a present sequence represents a human GEM content and a glioma cells. (M3) is useful for inducing an immune response to a glioma in a present sequence represents a human GEM content and a glioma cells.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                          2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 4e+02; Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hepatitis B virus (HBV) RNA target sequence #1053.
                                                                                                                                                                                                                                                                                                                                                                                   Sequence 17 BP; 1 A; 8 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           177 TGICAGCTGCTGGCCC 192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADM58919 standard; RNA; 17 BP.
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07-FEB-1994; 94US-00193227.
8-NOV-1999; 9US-00436430.
20-MAR-2000; 2000US-00531025.
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24-OCT-2000; 2000US-00696347.
08-JUN-2001; 2001US-00877478.
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity 87.5
les 14; Conservative
                                                                                                                                                                                                                                                                                                                                              invention.
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Novel enzymatic nucleic acid molecule such as DNAzymes and inozymes specifically cleaving RNA derived from hepatitis B virus and comprising one or more binding arms, useful for treating hepatitis and cirrhosis.

Morrissey D;

Mcswiggen JA,

Blatt L,

Draper K,

WPI; 2004-247781/23.

BLATT L. MCSWIGGEN J A. MORRISSEY D.

DRAPER K.

(DRAP/) (BLAT/) (MCSW/) MORR/)

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The invention relates to an enzymatic nucleic acid molecule that comprising one or more binding arms, without requiring the presence of a 2.-OH group within the molecule for activity. The nucleic acids are useful for treating hepatitis B virus infection, hepatitis, hepaticallular carcinoma, cirrhosis and liver failure, either alone or in combination with other therapies such as lamivudine and interferons. The mucleic acids are useful as diagnostic tools to examine genetic drift and mutations within diseased cells, for detecting the presence of HBV RNA in a cell, for the study of RNA and for down-regulating gene expression of a carget genes in bacterial, fungal, viral, plant or mammalian cells. This sequence represents an HBV RNA target sequence, used in the scope of the invention. Note: The sequence data for this patent is also available in electronic format from USPTO at sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Enzymatic nucleic acid molecules which specifically cleave RNA derived from hepatitis C virus (\mathrm{HCV}), useful for the treatment of \mathrm{HCV} infections,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to an enzymatic nucleic acid molecule which specifically cleaves RNA derived from hepatitis C virus (HCV), in which the binding arms of the enzymatic nucleic acid molecule comprises sequences complementary to any of the defined substrate sequences given in the specification. The nucleic acid molecule may be administered for the treatment of HCV infections, especially in combination with type I
                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; SEQ ID NO 917; 198pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HCV DNAzyme substrate sequence #917.
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                                                                                                                                                                                                                                                                                                                                                                                                            117 AGCGGGGGGAAAAGCC 132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADI83671 standard; RNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-DEC-2000; 2000US-00740332.
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                                                                                                                                                                                                                                                                                                                                                                                                                                     16 AGCGGCGGTAGAGCC
                                                                                                                                                                                                                                                                                                                                                     Local Similarity 87.5
les 14; Conservative
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MCSWIGGEN J.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ROBERTS E. PAVCO P A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hepatitis C virus.
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(PAVC/)
(MACE/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 473
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Matches

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from hepatitis C virus (HCV), useful for the treatment of HCV infections, especially in combination with type I interferon therapy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Snzymatic nucleic acid molecules which specifically cleave RNA derived
                                                                                                                                        enzymatic nucleic acid, RNA cleavage, hepatitis C virus, HCV, infection, type I interferon, DNAzyme.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ss; enzymatic nucleic acid; RNA cleavage; hepatitis C virus; HCV;
HCV infection; type I interferon; DNAzyme.
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                                                                                                                                                                                                                                                                                                                                                                                                                          Pavco PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; SEQ ID NO 1425; 198pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HCV DNAzyme substrate sequence #3755.
                                                                                                                 substrate sequence #1425
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                              ADI84179 standard; RNA; 17
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                                                                                    (first entry)
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                                                                                                                                                                                                                                                                                                                                                (MCSW/) MCSWIGGEN J.
(ROBE/) ROBERTS E.
(PAVC/) PAVCO P A.
(MACE/) MACEJACK D.
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                                                                                                                                                                                       Hepatitis C virus
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                                                                                                                 HCV DNAzyme
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            ADI84179
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Enzymatic nucleic acid molecules which specifically cleave RNA derived from hepatitis C virus (HCV), useful for the treatment of HCV infections, especially in combination with type I interferon therapy.
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                                                                                                   Gaps
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interferons. The present sequence represents a HCV DNAzyme substrate
                                                                                                                                                                                                                                                                                                                                                 enzymatic nucleic acid; RNA cleavage; hepatitis C virus; HCV; infection; type I interferon; DNAzyme.
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llarity 87.5%; Pred. No. 4e+02;
Conservative 0; Mismatches 2; Indels
                                                                    2.8%; Score 12.8; DB 1; Length 17; 75.0%; Pred. No. 4e+02; tive 2; Mismatches 2; Indels
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                                          Sequence 17 BP; 1 A; 9 C; 5 G; 0 T; 2 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; SEQ ID NO 3639; 198pp; English.
                                                                                                                                                                                                                                                                                                                      HCV DNAzyme substrate sequence #3639.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     223
                                                                                                                             210 ACCTGCGGCGGGTCGC 225
                                                                                                                                                                                                                                 ADI86393 standard; RNA; 17 BP
                                                                                                                                               ACCUGCGCCGCUCGC 17
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                                                                                                   Conservative
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MCSWIGGEN J.
ROBERTS E.
PAVCO P A.
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nes 14; Conserv
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                                                                                    Local Similarity
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(ROBE/) I
(PAVC/) I
(MACE/) R
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                                                                                                                                                                          Enzymatic nucleic acid molecules which specifically cleave RNA derived from hepatitis C virus (HCV), useful for the treatment of HCV infections, especially in combination with type I interferon therapy.
                                                                                                                                                                                                                               The invention relates to an enzymatic nucleic acid molecule which specifically cleaves RNA derived from hepatitis C virus (HCV), in which the binding arms of the enzymatic nucleic acid molecule comprises sequences complementary to any of the defined substrate sequences given in the specification. The nucleic acid molecule may be administered for the treatment of HCV infections, especially in combination with type i interferons. The present sequence represents a HCV DNAzyme substrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mutant cell identification-related mutagenic oligonucleotide SeqID752.
                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cell identification; oligonucleotide-directed sequence alteration; selectable phenotype; transgenic plant; herbicide resistance; sterile plant; abiotic stress tolerance; albino plant; amino acid production; ss.
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                                                                                                                                                                                                                                                                                                                                      / Match 2.8%; Score 12.8; DB 1; Length 17; Local Similarity 87.5%; Pred. No. 4e+02; les 14; Conservative 0; Mismatches 2; Indels
                                                                                                                                           Macejack D;
                                                                                                                                                                                                                                                                                                                     Sequence 17 BP; 1 A; 6 C; 6 G; 0 T; 4 U; 0 Other;
                                                                                                                                         Pavco PA,
                                                                                                                                                                                                              Claim 1; SEQ ID NO 3755; 198pp; English.
                                                                                                                                         Roberts E,
                                                                                                                                                                                                                                                                                                                                                                         433 GGACTCGCCTCACACA 448
                                            .8-DEC-2000; 2000US-00740332
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2003US-0453360P.
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                                                                                 BLATT L.
MCSWIGGEN J.
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                                                                                                   ROBERTS E.
PAVCO P A.
                                                                                                                      MACEJACK D
        US2003125270-A1.
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07-MAR-2003;
                           03-JUL-2003
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(ROBE/) 1
(PAVC/) 1
(MACE/) N
                                                                                                                                        Blatt L,
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This invention relates to a novel method of identifying a cell having a desired oligonucleotide-directed sequence alteration at a first nucleic acid target site within the cell. The method comprises identifying the desired sequence alteration in cells that have been selected for the presence of a selectable phenotype conferred by a concurrent oligonucleotide-directed sequence alteration at a second nucleic acid target site within the cells. The method is useful in identifying a cell having a desired oligonucleotide-directed sequence alteration at a first the production of plants with herbicide resistance, male or female sterile plants, abiotic stress tolerance, albino plants or plants with an altered amino acid production as well as for use in mammalian cell lines.
                                                                                                                                                                                                    Identifying a cell with a desired oligonucleotide-directed sequence alteration at a nucleic acid target site within the cell by identifying the desired sequence alteration in cells selected for the presence of a selectable phenotype.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence is that of a mutagenic oligonucleotide which was used in the exemplification of the invention.
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0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 17 BP; 3 A; 7 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                        Example 24; SEQ ID NO 752; 303pp; English.
(NAPR-) NAPRO BIO THERAPEUTICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (UYDE ) UNIV DELAWARE.
(NAPR-) NAPRO BIO THERAPEUTICS INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ilarity 87.5%;
Conservative
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                                                                   Kmiec EB, Van Brabant
                                                                                                                                    WPI; 2004-340941/31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-340941/31.
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es 14; Conserv
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Synthetic.
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This invention relates to a novel method of identifying a cell having a desired oligonuclectide-directed sequence alteration at a first nucleic acid target site within the cell. The method comprises identifying the desired sequence alteration in cells that have been selected for the presence of a selectable phenotype conferred by a concurrent oligonuclectide-directed sequence alteration at a second nucleic acid target site within the cells. The method is useful in identifying a cell having a desired oligonuclectide-directed sequence alteration at a first nucleic acid target site within the cell. The method may be useful for the production of plants with harbicide resistance, male or female sterile plants, abjoict stress tolerance, albino plants or plants with alterance, albino plants or plants with alterance amino acid production as well as for use in mammalian cell lines. The present sequence is that of a mutagenic oligonucleotide which was the sequence is that of a mutagenic oligonucleotide which was
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Identifying a cell with a desired oligonucleotide-directed sequence alteration at a nucleic acid target site within the cell by identifying the desired sequence alteration in cells selected for the presence of a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Anorectic; Antidiabetic; Muscular; Gene Therapy; CpG island; IGF2 gene intron 3; muscle mass; fat deposition; teat number; obesity; muscle deficiency; diabetes; SNP; single nucleotide polymorphism; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 4e+02; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 17 BP; 2 A; 5 C; 7 G; 3 T; 0 U; 0 Other;
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                                                                                            Example 24; SEQ ID NO 751; 303pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
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ADQ80740 standard; DNA; 17 BP.
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Matches 14; Conservative
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                                                            selectable phenotype
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(GENT-) GENTEC BV.
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The present invention relates to a method (M1) for selecting an animal for having desired genotypic or potential phenotypic properties. (M1) comprises testing the animal for the presence of a nucleic acid modification affecting: the activity of an evolutionary conserved CpG island located in intron 3 of an IGF2 gene; and/or binding of a nuclear factor to an IGF2 gene. The nuclear factor is capable of binding to a stretch of nucleotides which in the wild type pig, mouse or human IGF2 gene is part of an evolutionary conserved CpG island, located in intron 3 of the IGF2 gene. The stretch is functionally equivalent to (ADQ80709). The nucleic acid modification in ADQ80709 comprises a G to A translition at IGF2-intron3-nt3072. (M1) is useful for selecting an animal with properties related to muscle mass, fat deposition, and/or teat number. CC Also claimed is a method (M2) for modulating mNNA transcription of an IGF2 gene and/or modulating binding of a nuclear factor to an IGF2 gene. Also claimed is a method (M3) for identifying a compound capable of modulating mNNA transcription of an IGF2 gene and/or modulating of a nuclear factor to an IGF2 gene and/or modulating of a nuclear factor to an IGF2 gene and/or modulating planding of a nuclear factor to an IGF2 gene and/or modulating binding of a nuclear factor to an IGF2 gene and/or modulating binding of a nuclear factor to an IGF2 gene and/or modulating binding of a nuclear factor to an IGF2 gene and/or modulating binding of a nuclear factor to an IGF2 gene and/or modulating binding of a nuclear factor to an IGF2 gene and/or modulating binding of a nuclear factor to an IGF2 gene and/or modulating binding of a nuclear factor to an IGF2 gene and/or modulating binding of a nuclear factor to an IGF2 gene and/or modulating binding of a nuclear factor to an IGF2 gene and/or modulating an activity muscle deficiencies and diabetes. The present sequence is a porcine sequence tagged sites (STS) comprising a porcine sequence in an example from the invention.
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androgen signal transduction pathway protein;
androgen signal transduction; androgen receptor; hair loss;
hyperandrogenic condition; androgenic alopecia; male pattern alopecia;
acne vulgaris; seborrhoea; female hirsutism; prostatic hypertrophy; ds
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/note= "2 deoxynucleotide overhang"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 17 BP; 0 A; 14 C; 2 G; 1 T; 0 U; 0 Other;
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                      3xample 1, Page 20, 38pp, English.
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Best Local Similarity 87.5
Matches 14; Conservative
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The present invention relates to novel small interfering RNAs (siRNAs), comprising a double stranded RNA, where one strand comprises a partial nucleic acid sequence of an anadrogen signal transduction pathway protein, and where the double-stranded RNA inhibits translation of mRNA encoding the nucleic acid sequence of the androgen signal transduction pathway. The compress blocking the androgen signal transduction pathway. The androgen signal transduction pathway. The androgen signal transduction pathway is captured to a comparable (ADQ92455 and ADQ92516), the androgen are (ADQ92571), aromatase (ADQ92455 and ADQ92516), the androgen and II of 5-alpha reductase (ADQ93182), 3-beta-chydroxysteroiddehydrogenase (ADQ93182), 3-beta-chydroxysteroidde
                                                                                                                                                Inhibitory nucleic acid that inhibits expression of an androgen signal transduction pathway protein useful for treating hair loss, comprises a double stranded RNA having a partial sequence encoding a pathway protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; gene therapy; thymidine kinase gene; anticancer therapy; human; mutagenesis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human telomerase RNA gene (hTR) promoter specific primer hllc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 4e+02; 1ve 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                Claim 11; Page 41; 92pp; English.
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Best Local Similarity 87.5
Matches 14; Conservative
                                                                                                  WPI; 2004-561892/54.
  (GENC-) GENCIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the invention
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                                                     Kahn S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 481
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     8
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The invention relates to promoter regions from mouse and human telomerase RNA (TR) component genes. The TR gene promoter can be linked to a heterologous gene, especially a gene encoding a cyctoxin, for therapy of cancer, especially neoplasias. The telomerase is necessary for the concer, especially neoplasias. The telomerase is necessary for the concer, especially neoplasias. The telomerase is necessary for the concertiated proliferative capacity of many human cancers. Mutation or concertiated proliferative capacity of many human cancers. Mutation or concertiated proliferative can be used to block transcription from the TR gene promoter through interaction of the 5' regulatory sequences. These canbstances, e.g. antisense oligomucleotides, transcription factors. Oppide mucleic acids and factors that disrupt signal transduction, are seful for cancer therapy. In particular, gene therapy vectors (especially pdf62-codaups) comprising the promoter and a viral thymidine concertaints and be used to convert a prodrug, e.g. gancyclovir, so that neoplasia can be controlled or treated. Direct down-regulation of telomerase RNA gene through manipulation of transcription factors may be effective anticancer therapy and the cloning of the hTR gene promoter callows the analysis of therapeutic molecules which modulate hTR promoter and manipulation of activity, Sequences Assorbede S121 represent PCR primers used in cloning
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                            Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RNA component; human telomerase; antisense oligonucleotide; infection; neuroblastoma; bladder cancer; colon cancer; prostate cancer; cancer; contraception; sterilisation; immunosuppression; therapeutic; hTR; immune system down-regulation; anti-inflammatory therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 2.8%; Score 12.8; DB 1; Length 25; Best Local Similarity 70.8%; Pred. No. 5.3e+02; Matches 17; Conservative 0; Mismatches 7; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RNA component of human telomerase (hTR) antisense oligo 14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        and mutagenisis of human TR gene (hTR) promoter region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Wu F, Kealey JT, Pruzan R, Weinrich SL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 25 BP; 1 A; 5 C; 15 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         223 CGCCTGCCCAGCCCCGAACCCCG 246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25 CCCAGGCCCACCTCCGCACCCG 2
                                                                                                                                              Disclosure, Fig 12, 109pp, English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              97WO-US023619.
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96US-00770565.
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                                       WPI; 1999-479183/40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GERO-) GERON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         08-OCT-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-DEC-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    02-JUL-1998
  Keith WN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAV41169;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
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                                                                                                               Sequences shown in AAV41169 to AAV41181 represent antisense
oligonucleotides to the RNA component of human telomerase (hTR). These
antisense oligonucleotides specifically hybridise to a nucleotide
equence within an accessible region of the hTR, but that does not
oligonucleotides may specifically be used for detection of an RNA
component of human telomerase in a sample. This is useful for diagnosing
cancer (especially neuroblastoma, bladder, colon and prostate cancer),
and providing prognosis for a cancer patient. The inhibitory
cancer (especially neuroblastoma, bladder, colon and prostate cancer),
and providing prognosis for a cancer patient. The inhibitory
cancer (especially neuroblastoma, bladder, colon and prostate cancer),
and providing prognosis for a cancer patient. The inhibitory
cancer (especially neuroblastoma cancer patient. The inhibitory assembly of
interfering with transcription of the RNA component, decreasing the half-
coligonucleotides can inhibit the component transcript, inhibiting assembly of
the RNA component into the telomerase holoenzyme, or inhibiting telomerase activity in both cultured cells and in
be used for inhibiting telomerase. These antisense oligonucleotides can
be used for inhibiting telomerase activity in both cultured cells and in
cells in vivo. They can be used in therapeutics for treating or
immunosuppression, and for selectively down-regulating specific branches
of the immune system, e.g. a specific subset of T-cells, in anti-
inflammatory therapies or for treating infections by, e.g. yeast,
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                  - used for detecting
            New polynucleotide(s) anti:sense to human telomerase - used for detecting or inhibiting human telomerase, e.g. for treating cancers, contraception, immuno-suppression or treating infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; gene therapy; thymidine kinase gene; anticancer therapy; human; mutagenesis; PCR primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human telomerase RNA gene (hTR) promoter specific primer h112.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.8%; Score 12.8; DB 1; Length 30; 70.8%; Pred. No. 5.4e+02; tive 0; Mismatches 7; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 30 BP; 4 A; 10 C; 8 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RES CAMPAIGN TECHNOLOGY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       131 CCTCGGCCTGCCGCCTTCCACCGT 154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ccrcrrccrcccccraaacccr 28
                                                                                     Claim 11; Page 65; 80pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               298/c
AAZ07298 standard; DNA; 38 BP.
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Matches 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-JAN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAZ07298;
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gene therapy

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The intention retails to produce regions from mouse and intentions of the component genes. The TR gene promoter can be linked to a heterologous gene, especially a gene encoding a cytotoxin, for therapy of cancer, especially appeals are the telomerase is necessary for the unrestricted proliferative capacity of many human cancers. Mutation or dysregulation of the telomerase expression pathway may cause reactivation or dysregulation of telomerase expression in cancer. Substances, cor upregulation of telomerase expression in cancer. Substances, cor upregulation of telomerase expression in cancer. Substances, compared to the S' regulatory sequences. These gene promoter through interaction of the S' regulatory sequences. These causeful for cancer therapy. In particular, gene thrangulation, are useful for cancer therapy. In particular, gene thrangulation of especially pGT62-codAupp) comprising the promoter and a viral thymidine kinese gene can be used to convert a prodrug, e.g. gancyclovir, so that neoplasia can be controlled or tracated. Direct down-regulation of telomerase RNA gene through manipulation of transcription factors may be effective anticancer therapy and the cloning of the HTR gene promoter allows the analysis of therapeutic molecules which modulate HTR promoter activity. Sequences AAZO7696-321 represent PCR primers used in cloning and mutagenisis of human TR gene (HTR) promoter region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                      invention relates to promoter regions from mouse and human telomerase
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pair of RNA molecules for detecting telomerase, useful for diagnosis of cancer or metastases, can be ligated when bound to telomerase subunit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human, telomerase, hTR, telomeric repeat amplification protocol; TRAP,
identification, detection, quantification, cancer, metastasis, 8s.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 38 BP; 3 A; 7 C; 21 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Adenovirus nucleotide sequence SEQ ID NO:20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          223 CGCCTGCCCAGCCCCCGAACCCCG 246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CCCAGGCCCACCCTCCGCAACCCG 2
Disclosure, Fig 12, 109pp, English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 70.8°
Matches 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12-JUN-1997;
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= RNA segment of 10 to 100000 nucleotides (nt) that together are replicated by RNA replicase; B and E = RNA segments of 10 to 250 nt trom the Y region of human telomerase and bind specifically to (1); C and D = RNA segments of 10 to 250 nt from the Y region of human telomerase and bind specifically to (1); C and D = C and D produces R3 of formula 5.-A-B-C-D-E-F-3' (R3) with E and B bound to (1). Replication of R3 by RNA replicase indicates presence of (1) to (1). Replication of R3 by RNA replicase indicates presence of (2) to this process containing R1, R2, ligase and an amplification system; and (3) method for making R1 R2, ligase and an amplification system; and (3) method for detect and quantify telomerase, particularly for diagnosis of cancer and for detection of metastases. R1 and R2 provide an assay that does not require expensive equipment or highly trained personnel, and is suitable for automation. The present sequence invention
RNA segment of 10 to 100000 nucleotides (nt) that together
                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention
85999599999999
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Sequence 62 BP; 16 A; 27 C; 10 G; 9 T; 0 U; 0 Other; 2.8%;

Score 12.8; DB 1; Length 62; Pred. No. 3.2e+02; 0; Mismatches 7; Indels 0; Mismatches 222 TCGCCTGCCCAGCCCCGAACCCC 245 61 38 rcccagcccacccrccgcaaccc 70.8%; Query Match
Best Local Similarity 70.0.
Best Local 17; Conservative 쉽 à

0

Gaps

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485

AAH24816 standard; RNA; 62 GGGGGGGGX8X444X8X1X3X8X8X8X8X8X8X8X8X8X8X8X8X8X8

(first entry) 06-AUG-2001 AAH24816;

Human nucleic acid sequence derived from Y-1 domain of telomerase.

RNA-binding protein; RBP; RNA replicase; RNA identification; telomerase;

Homo sapiens.

US6238867-B1

29-MAY-2001

22-FEB-1999;

99US-00255464.

98US-0075495P.

(INVI-) INVITRO DIAGNOSTICS INC.

Grossman A; Roninson IB,

WPI; 2001-366472/38.

RNA New ribonucleic acids useful for identifying naturally occurring RNA sequences having affinity for RNA-binding protein having protein and Example 2; Col 26; 36pp; English. components.

The specification describes a first RNA (RNA1) and a second RNA (RNA2) for use in binding an RNA-binding protein (RBP) having protein and RNA components. RNA1 has the formula 5'-A-B-C-3', where A is section having 10-100,000 nucleotides and is can be received by an RNA replicase and with another DNA sequence, F, being replicated; B is section having 10-3,000 nucleotides which have affinity to one consensus sequence of the RBP and which can bind to the protein component; C is section having about 1-20 nucleotides and which can be ligated to D of the second RNA molecule. RNA2 has the formula S'D-B-F-3', where D is section having 1-20 nucleotides and which can be ligated to C; E is section 10-3,000 nucleotides which have affinity to second consensus sequence of the RBP

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The present invention describes a pair of RNA molecules (R1, R2) for detecting a first subunit protein (I) of telomerase. R1 and R2 both bind to (1) and have formulae 5'-A-B-C-3' (R1) 5'-D-B-F'3' (R2) where: A and F = RNA segment of 10 to 100000 nucleotides (nr) that together are replicated by RNA replicase; B and E = RNA segments of 10 to 250 nt from the Y region of human telomerase and bind specifically to (1); C and D = RNA segments of 1 to 10000 nt which can be ligated together. Ligation of C and D produces R3 of formula 5'-A-B-C-D-E-F-3' (R3) with E and B bound C and D produces R3 of formula 5'-A-B-C-D-E-F-3' (R3) with E and B bound C C and D produces R3 of formula 5'-A-B-C-D-E-F-3' (R3) with E and B bound C C and D produces R3 of formula 8'-A-B-C-D-E-F-3' (R3) with E and B bound C C and (3) method for detecting (I) using R1 and R2' (1) kit for this process containing R1, R2, ligase and an amplification system; and (3) method for making R1 and R2 by transcription from appropriate DNA. R1 and R2 are used to detect and quantify telomerase, particularly for diagnosis of cancer and for detection of metastases. R1 and R2 process contained and assay that does not require expensive equipment or highly trained personnel, and is suitable for automation. The present sequence represents an oligonucleotide used in the exemplification of the present
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and which can bind to the protein component; F is section having 10-100,000 nucleotides which can be received by an RNA replicase and with A being replicated. RNA1 and RNA2 are capable of forming a third RNA (RNA3) of formula 5'A-B-C-D-E-F'3'. The method is useful for the identification and characterization of RNA sequences having specific affinity to amino acid consensus sequences of RBPs, and to RNAs. AAH24815-16 were used to produce a double-stranded RNA1, comprising the Y-1 domain of human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; telomerase; hTR; telomeric repeat amplification protocol; TRAP; identification; detection; quantification; cancer; metastasis; ss.
                                                                                                                                                                                                                                                                                   Gaps
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0
                                                                                                                                                                                                                                   2.8%; Score 12.8; DB 1; Length 62; 70.8%; Pred. No. 3.2e+02;
                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                         Sequence 62 BP; 16 A; 27 C; 10 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Adenovirus nucleotide sequence SEQ ID NO:19.
                                                                                                                                                                                                                                                         Pred. No. 3.2e
0; Mismatches
                                                                                                                                                                                                                                                                                                                  222 TCGCCTGCCCAGCCCCCGAACCCC 245
                                                                                                                                                                                                                                                                                                                                                           rcccaggcccacccrccgcaaccc 61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAA08204 standard; DNA; 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-282223/24.
                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mastadenovirus.
                                                                                                                                                                                                                                   Mac
Local Sim
17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US6037126-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-MAR-2000
                                                                                                                                                      telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Grossman A;
                                                                                                                                                                                                                                                                                                                                                              38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAA08204;
                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein.
                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 486
                                                                                                                                                                                                                                                                          Matches
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                     용
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The specification describes a first RNA (RNA1) and a second RNA (RNA2) for use in binding an RNA-binding protein (RBP) having protein and RNA components. RNA1 has the formula 5'-A=C-3', where A is section having 10-100,000 nucleotides and is can be received by an RNA replicase and with another DNA sequence, F, being replicated; B is section having 10-3,000 nucleotides which have affinity to one consensus sequence of the RNB and which can bind to the protein component; C is section having 10-300 nucleotides and which can be ligated to D of the second RNA molecule. RNA2 has the formula 5'-D-E-F-3', where D is section having 1-30 nucleotides which have affinity to second consensus sequence of the RNB continue and which can be ligated to C; E is section 10-3,000 nucleotides which have affinity to second consensus sequence of the RNB continue and protein component; F is section having 10-10,000 nucleotides which can be received by a RNA replicase and with A being replicated. RNA1 and RNA2 are capable of forming a third RNA (RNA3) of formula 5'A-B-C-D-E-F-3'. The method is useful for the identification and characterization of RNA sequences having specific affinity to amino acid consensus sequences of RRB9, and to RNAs. AAH24815-16 were used to produce a double-stranded RNA1, comprising the Y-1 domain of human
                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New ribonucleic acids useful for identifying naturally occurring RNA sequences having affinity for RNA-binding protein having protein and RNA
                                                                                                                                                                                                                                                                                                    RNA-binding protein; RBP; RNA replicase; RNA identification; telomerase;
                                                                 Gaps
                                                                                                                                                                                                                                                                         Human nucleic acid sequence derived from Y-1 domain of telomerase.
                                                                 ö
                                    Score 12.8; DB 1; Length 66;
Pred. No. 3e+02;
0; Mismatches 7; Indels
                                                              7; Indels
              28 G; 17 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Seguence 66 BP; 10 A; 11 C; 28 G; 17 T; 0 U; 0 Other;
                                                                                            222 TCGCCTGCCCAGCCCCGAACCCC 245
                                                                                                                    29 receasseceácerecisaaces 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 2; Col 26; 36pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (INVI-) INVITRO DIAGNOSTICS INC.
                                                                                                                                                                                         BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                      98US-0075495P
                                                                                                                                                                                                                                                                                                                                                                                                                            99US-00255464
                                    Query Match 2.8%;
Best Local Similarity 70.8%;
Matches 17; Conservative
            Sequence 66 BP; 10 A; 11 C;
                                                                                                                                                                                         AAH24815 standard; RNA; 66
                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Grossman A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-366472/38.
                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                            22-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Roninson IB,
                                                                                                                                                                                                                                                                                                                                                                     US6238867-B1
                                                                                                                                                                                                                                               06-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                 29-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            components.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                telomerase
                                                                                                                                                                                                                    AAH24815
                                                                                                                                                             RESULT 487
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The invention comprises the amino acid and coding sequences of the human cholesteryl ester transfer protein (CETP), the invention also comprises bolymorphisms identified within the CETP gene. The DNA and protein sequences of the invention are useful in haplotyping and/or genotyping the CETP gene in an individual. The DNA and protein sequences may also be used to screen drugs or compounds targeting the CETP or its variant to treat a condition or disease associated with CETP (e.g. atherosclerosis, cardiovascular disease or hypercholesterolaemia). The present DNA sequence represents an allele specific oligonucleotide PCR primer for the
                                                                                                                                                                                                                                                                                                                                                                   human; cholesteryl ester transfer protein; CETP;
single nucleotide polymorphism; SNP; drug screening; atherosclerosis;
cardiovascular disease; hypercholesterolasemia;
allele specific oligonucleotide; ss; PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New isolated polynucleotide useful for haplotyping and/or genotyping cholesteryl ester transfer protein (CETP) gene in an individual or in screening for drugs useful in treating diseases associated with CETP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                 Human CETP gene allele specific oligonucleotide PCR primer #21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Lachowicz M,
Tirrell C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 43; SEQ ID NO 80; 250pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          completed: August 24, 2005, 14:24:50
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Sausker EA,
TCCCAGGCCCACCCTCCGCAACCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GENA-) GENAISSANCE PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-APR-2003; 2003WO-US013288.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-APR-2002; 2002US-0375791P.
                                                                                                                                                      15
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                                                                                                                                                                                                                                                             (first entry)
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                                                                                                                                                   ADG98448 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ĄΞ
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N, Rounds I
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es 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          human CETP gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO2003091277-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Petersen N,
                                                                                                                                                                                                                                                             11-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     36-NOV-2003
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                   29
                                                                                                                                                                                                          ADG98448;
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                                                                                                    RESULT 488
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Gaps

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2.8%; Score 12.8; DB 1; Length 66; 70.8%; Pred. No. 3e+02; ive 0; Mismatches 7; Indels

Query Match
Best Local Similarity 70.8
Matches 17; Conservative

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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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- nucleic search, using sw model OM nucleic

August 24, 2005, 14:51:46; Search time 0.001 Seconds (without alignments) 17.138 Million cell updates/sec ö

Run

US-09-436-060A-16

Perfect score:

1 gggttgcggagggtgggcct.....aggactcggctcacacatgc 451 Sequence:

IDENTITY_NUC Gapop 10.0 , Gapext 0.5 Scoring table:

1 segs, 19 residues Searched:

a Total number of hits satisfying chosen parameters:

seq length: 0 seq length: 200000000 Minimum DB Maximum DB

Post-processing: Minimum Match 10% Maximum Match 100% Listing first 2 summaries

rst.subdb:* Database : Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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> ACCESSION: AA934303 ACCESSION: AA934303 1 AA934303 1 AA934303 Query Score Match Length DB 19 8 9 7 0 14.4 8.8 Result Š

ALIGNMENTS

AA934303
SWO-LICANI2H12 Onchocerca volvulus infective larva cDNA
(SAM94WL-OvL3) Onchocerca volvulus cDNA clone onch672 5' similar to
TR:033571 033571 ATPASE SUBUNIT 6 ;, mRNA sequence. Eukaryota, Metazoa, Nematoda, Chromadorea, Spirurida, Filarioidea, Onchocercidae, Onchocerca.

Onchocercidae, Onchocerca.

I (Dases I to 19)

Williams, S.A., Lizotte-Waniewski, M., Laney, S., Wenhong, L.,
Hillar, L., Allen, M., Bowles, L., Geisel, S., Jost, S., Kucaba, T.,
Martin, J., Steptce, M., Theising, B., White, Y., Wylie, T.,
Chappell, J., Person, B., Gibbons, M., Harvey, N., Pape, D.,
Chamberlain, A., Morales, R., Schurk, R., Ritter, E., Kohn, S.,
Underwood, K. and Marra, M. Molecular Parasitology Smith College Department of Biological Sciences Department of Biological Sciences, Clark Science Center, Smith Contact: Steven A. Williams AA934303.1 GI:3091460 Onchocerca volvulus Unpublished (1998) AA934303 VERSION KEYWORDS SOURCE ORGANISM DEFINITION TITLE JOURNAL COMMENT ACCESSION REFERENCE AUTHORS RESULT 1 AA934303

Email: genome@smith.edu
The library was constructed by Wenhong Lu. The library is available
from Dr. S.A. Williams, email genome@smith.edu When requesting this
clone from Dr. Williams, please reference the Williams lab clone id

Fax: 4135853786

Unpublished (1998)
Contact: Steven A. Williams
Molecular Parasitology
Smith College Department of Biological Sciences
Department of Biological Sciences, Clark Science Center, Smith
College, Northampton, MA, 01063, USA

TITLE JOURNAL

COMMENT

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Email: genome@smith.edu
The library was constructed by Wenhong Lu. The library is available from Dr. S.A. Williams, email genome@smith.edu When requesting this oflone from Dr. Williams, please reference the Williams lab clone id SWOYL3CANI2H12
SWOYL3CANI2H12
Seq primer: -40ml3 fwd. ET from Amersham
High quality sequence stop: 1.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Anotes "Vector: lambda UniZap XR; Site 1: EcoR I; Site 2: Xho I; Cutaneous filarial nematode parasite of humans. The I; Cutaneous filarial nematode parasite of humans. The I construction of control isolated from mosquitoes IO days after infection and converted to double stranded cDNA using reverse transcriptase and oligo (dT) followed by RNase H and DNApol I. The library had 1.8 x 10E5 independent recombinants and average insert size was 900 base pairs. The library was constructed by Wenhong Lu. The library is available from Dr. S.A. Williams, email genome@smith.edu."
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SWO-KISCANI2H12 Onchocarca volvulus infective larva cDNA
(SAM94ML-OvL3) Onchocerca volvulus cDNA clone onch672 5' similar to
TR:Q33571 Q33571 ATPASE SUBUNIT 6 ;, mRNA sequence.
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Eukaryota; Merazoa; Nematoda; Chromadorea; Spirurida; Filarioidea;
Onchocercida; Onchocerca.

1 (bases 1 to 19)
Williams,S.A., Lizotte-Waniewski,M., Laney,S., Wenhong,L.,
Williams,S.A., Lizotte-Waniewski,M., Geisel,S., Jost,S., Kucaba,T.,
Martin,J., Steptoe,M., Theising,B., White,Y., Wylie,T.,
Chappell,J., Person,B., Gibbons,M., Harvey,N., Pape,D.,
Chamberlain,A., Worales,R., Schurk,R., Ritter,E., Kohn,S.,
Molecular Parasitology OvL3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /clone="onch672"
/lab host="XIJ-Blue MRF'"
/lab host="XIJ-Blue MRF'"
/clone lib="Onchocerca volvulus infective larva cDNA
(SAW94WI-OvL3)"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3.2%; Score 14.4; DB 1; Length 19; 88.2%; Pred. No. 0; tive 0; Mismatches 2; Indels

    19
    organism="Onchocerca volvulus"

Northampton, MA, 01063, USA
                                                                                                                                                                                                                                                                                                                                                                                    /mol_type="mRNA"
/strain="Sierra Leone"
/db_xref="taxon:6282"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  162 AGAGCAAACAAAATG 178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AGAGNAAACAAAAAAG 17
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                                                             Fax: 4135853786
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Best Local Similarity
....hes 15; Conserve
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AA934303/c
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KEYWORDS
                                                                                                                                                                                                                                                                                                 FEATURES
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- SWOVLJCAN12H12
Seq primer: -40ml3 fwd. ET from Amersham
High quality sequence stop: 1.

Location/Qualifiers

1. .19
| organism="Onchocerca volvulus" |
| find train="Sierra Leone" |
| strain="Sierra Leone" |
| strain="Sierra Leone" |
| strain="Sierra Leone" |
| clone="Inb="Nonchocerca volvulus infective larva cDNA (SAM94ML-OVL) |
| clone="Inb="Nonchocerca volvulus infective larva cDNA (SAM94ML-OVL) |
| sam94ML-OVL) |
| clone="Inb="Nonchocerca volvulus infective larva of from kNNA was prepared from third stage infective larvae of of Onchocerca volvulus isolated from mosquitoes 10 days after infection and converted to double stranded cDNA using reverse transcriptase and oligo(dT) followed by RNAse H and DNApol I: The library had 1.8 x 10E5 independent recombinants and average insert size was 900 base pairs. The library was constructed by Wenhong Lu. The library is available from Dr. S.A. Williams, email genome@smith.edu."
                                                                                                                              FEATURES
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; 0 Gaps . DB 1; Length 19; Query Match 2.0%; Score 8.8; DB 1; Length 19 Best Local Similarity 83.3%; Pred. No. 0; Matches 10; Conservative 0; Mismatches 2; Indels

36 CATTTTTTGTCT 47

ò 임 Search completed: August 24, 2005, 14:51:46 Job time : 0.001 secs

2234336	000000000000000000000000000000000000000		sequence 36, Appl Sequence 40, Appl Sequence 28, Appl Sequence 29, Appl Sequence 2, Appl Sequence 2, Appl Sequence 2, Appl Sequence 25, Appl Sequence 26, Appl Sequence 26, Appl Sequence 26, Appl Sequence 27, Appl Sequence 27, Appl Sequence 21, Appl Sequence 27, Appl Sequence 28, Appl Sequence 2, Appl Seq
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US-10-330-872-2 US-08-330-123A-25 US-08-482-115B-25 US-08-660-678A-25	US-08-485-778-33 US-08-472-802C-26	US-08-998-443-25	US-09-060-523-25 US-09-580-517-25	US-09-057-351-25 US-08-833-377-3	US-08-330-123A-4	US-08-330-123A-7	US-08-462-1132-7	US-08-660-678A-26	US-08-660-678A-29	US-08-770-565-6	US-08-770-565-7	US-08-770-565-13	US-U8-485-778-6 US-08-485-778-9	US-08-472-802C-8	US-08-472-802C-41	US-08-833-377-7	US-08-320-330A-8 US-08-520-550A-9	US-08-630-019A-13	US-08-838-545-41	US-08-998-443-7	US-U8-338-443-26 US-08-998-443-29	US-09-060-523-7	US-09-349-532-41	US-U9-58U-51/-4 US-09-580-517-7	US-09-057-351-7	US-09-057-351-40	US-08-770-565-9 US-08-838-545-60	US-09-349-532-60	US-09-018-125-2	US-U8-833-3//-14 US-08-838-545-9	US-09-349-532-9	US-08-770-565-14	US-U8-974-549A-543 US-09-402-181B-543	US-09-721-456-543	US-08-026-143B-13	PCT-US92-10621-13 BCT-11594-0223-13	US-08-838-545-27	US-09-349-532-27	US-09-345-882-76	US-09-596-938-11	US-08-770-303-10 US-08-630-019A-12	US-08-630-019A-18	US-08-630-019A-40	US-08-838-545-2	US-U8-838-343-3 IIS-08-838-545-28	US-08-838-545-45	US-09-349-532-2	US-09-349-532-5	US-09-349-532-28 TR-09-349-532-45	US-UY-34Y-53Z-45 US-09-673-298-4	US-UY-6/3-230-4
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; Sequence 15, Application US/09255464B
; Patent No. 6238867
PRIOR APPLICATION NUMBER: 60/6;
PRIOR FILING DATE: 1999-02-23;
NUMBER OF SEQ ID NOS: 25
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 16
LENGTH: 62
                                            1998-02-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 10570
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 100.0
Matches 54; Conservative
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                           ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear;
MOLECULE TYPE: DNA
US-08-873-709-19
                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 54; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20, Application US/08873709
Sequence 20, Application US/08873709
Setent No. 6037126
GENERAL INFORMATION:
APPLICANT: Grossman, Abraham
TITLE OF INVENTION: COMPOSITIONS, METHODS, KITS AND
TITLE OF INVENTION: PROTEIN COMPONENT OF TELOMERASE ENZYME
NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
ADDRESSE: Abraham Grossman
                                                 Sequence 212, App
Sequence 9, Appli
Sequence 106, App
Patent No. 5166057
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Sequence 16, Application US/09255464B

Sequence 10, Application US/09255464B

Sequence 10, Application US/09255464B

GENERAL INFORMATION:
APPLICANT: Roninson, Igor
TITLE OF INVENTION: Identifying Naturally Occurring RNA Sequences Having
TITLE OF INVENTION: Affinity for RNA-Binding Proteins
FILE REPERENCE: Q001/004B

FILE REPERENCE: Q001/004B

CURRENT APPLICATION NUMBER: US/09/255,464B
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        Sequence 9
Sequence 2
Sequence 9
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM FC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/873,709

FILING DATE: 12-JUN-1997

CLASSIFICATION: 435

ATTORNEY/AGENT INFORMATION:

NAME: Janiuk, Anthony J.

REGISTRATION NUMBER: 29,809

REGISTRATION NUMBER: 29,809

REGISTRATION NUMBER: 29,001

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION
        US-08-871-732A-9
US-09-180-437-212
US-09-346-510B-9
US-09-544-934B-106
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Pred. No. 0.74;
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                                                                                                                                                                                  5166057-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Pleasantville STATE: NY STATE:
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US-08-873-709-20
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US-08-873-709-20/c
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Best Local 8
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TITLE OF INVENTION: COMPOSITIONS, METHODS, KITS AND
TITLE OF INVENTION: APPRATUS FOR DETERMINING THE PRESENCE OR ABSENCE OF
TITLE OF INVENTION: APPRATUS FOR DETERMINING THE PRESENCE OF
TITLE OF INVENTION: PROTEIN COMPONENT OF TELOMERASE BNZYME
NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
ADDRESSEE: Abraham Grossman
STREET: 666 Washington Avenue
CITY: Pleasantville
CITY: Neasantville
                                                   Gapв
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gapa
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DB 1; Length 62; 0.74;
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COMPUTER: FIGHTY COMPATIBLE
COMPATER: PRICE COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/873,709
FILING DATE: 12-JUN-1997
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: ADAILY, Anthony J.
REGISTRATION NUMBER: 29,809
REPRENCE/POCKET NUMBER: 29,809
REPRENCE/POCKET NUMBER: 29,809
REPRENCE/POCKET NUMBER: 29,809
REPRENCE/POCKET NUMBER: 29,809
RELECOMMUNICATION INFORMATION:
TELECHOMENICATION FOR SEQ ID NO: 19:
SEQUENCE CHARACTERISTICS:
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0.81;
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  12.0%; Score 54; DB
100.0%; Pred. No. 0.7
:ive 0; Mismatches
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Patent No. 6037126
GENERAL INFORMATION:
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MOLECULE TYPE: RNA (genomic)
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US-09-717-828B-1/c
              US-08-838-545-49
                                                                                                                                                                                                                                                                                  RESULT 6
US-09-349-532-49
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
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            APPLICANT: Roninson, Igor
APPLICANT: Grossman, Abraham
TITLE OF INVENTION: Compositions, Methods, Kits and Apparatus for
TITLE OF INVENTION: Compositions, Methods, Kits and Apparatus for
TITLE OF INVENTION: Identifying Naturally Occurring RNA Sequences Having
TITLE OF INVENTION: Affinity for RNA-Binding Proteins
TITLE OF INVENTION: Affinity for RNA-Binding Proteins
TITLE OF INVENTION WIMBER: US/09/255,464B
CURRENT APPLICATION NUMBER: 60/075,495
PRIOR PRILING DATE: 1998-02-23
NUMBER OF SEQ ID NOS: 25
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 15
LENGTH: 66
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APPLICANT: Wright, Woodring E.
APPLICANT: Wright, Woodring E.
APPLICANT: Wright, Weczyslaw A.
APPLICANT: Corey, David R.
APPLICANT: Corey, David R.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: 60 CORRESPONDENCE ADDRESS: 60 CORRESPONDENCE ADDRESS:
ADDRESSER: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 54; DB 1; Length 66;
Pred. No. 0.81;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
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SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
CLASSIFICATION NUMBER: US/08/630,019
FILING APPLICATION DATA:
APPLICATION NUMBER: US 08/630,019
FILING APPLICATION NUMBER: US 08/630,019
FILING APPLICATION NUMBER: US 08/630,019
FILING APPLICATION NUMBER: 32,944
REPERENCE/DOCKET NUMBER: 015389-001610US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFRENCE: (415) 576-0200
INFORMATION FOR SEQ ID NO: 49:
SEQUIENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GGGTTGCCGGAGGGTGGGCCTGGGAGGGGTGGTGGCCATTT
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100.0%; Pred. No. v...
... 0; Mismatches
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 49, Application US/08838545
Patent No. 6046307
GENERAL INFORMATION:
APPLICANT: Shay, Jerry W.
                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 100."
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STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA ORGANISM: Homo sapiens
  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       USA
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APPLICANT: Shay, Jerry W.

APPLICANT: Wight, Woodring E.

APPLICANT: Platyszek, Mieczyslaw A.

APPLICANT: Ozrey, David R.

APPLICANT: No. 6294650ton, James C.

TITLE OF INVENTION: Modulation of Mammalian Telomerase by
TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60

CORRESPONDENCE ADDRESS:
  DB 1; Length 31;
                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ch (5.9%; Score 31; DB 1; Length 31; 1 Similarity 71.0%; Pred. No. 20; 22; Conservative 9; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/349,532
                                               ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Townsend and Townsend and Crew LLP STREEF: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California
score 31; DB 1; Pred. No. 20; 9; Mismatches
                                                                                          40 TITIGICIAACCCIAACTGAGAAGGGCGIAG 70
                                                                                                                    1 UUUUGUCUAACCCUAACUGAGAAGGGCGUAG 31
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CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/838,545
FILING DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                     Sequence 49, Application US/09349532
Patent No. 6294650
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Storella, John R. REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 01:
TELECOMMUNICATION: TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: RNA (genomic)
Query Match 6.9%;
Best Local Similarity 71.0%;
Matches 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 31 base pairs
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NAME/KEY: misc_feature LOCATION: (31)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LOCATION: (1) —
OTHER INFORMATION: Biotin 5'-terminal
NAME/KEY: misc_feature
LOCATION: (31)
OTHER INFORMATION: Biotin 3'-terminal
OTHER INFORMATION: Description of Artificial Sequence: Affinity Agent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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GENERAL INFORMATION:
APPLICANT: Weinrich, Scott L
APPLICANT: Atkinson III, Edward M
APPLICANT: Lichtsteiner; Serge P
APPLICANT: Lichtsteiner; Serge P
APPLICANT: Lichtsteiner; Serge P
APPLICANT: Pruzan, Ronald A
TITUE OF INVENTION: A Method for Purifying Telomerase
CURRENT FILICATION NUMBER: US/09/71,829A
CURRENT FILING DATE: 2000-11-20
PRIOR APPLICATION NUMBER: 09/420,056
PRIOR FILING DATE: 1999-10-18
PRIOR FILING DATE: 1999-10-18
PRIOR RELIGHTON NUMBER: 08/833,377
PRIOR APPLICATION NUMBER: 08/813,377
PRIOR APPLICATION NUMBER: 08/510,736
PRIOR FILING DATE: 1995-08-04
NUMBER OF SEQ ID NOS: 11
SOFTWARE: PatentIn Ver. 2.1 edited
SEQ ID NO 1
LENGTH: 31
                                                                 APPLICANT: Weinrich, Scott L
APPLICANT: Atkinson III, Edward M
APPLICANT: Atkinson III, Edward M
APPLICANT: Lichtsteiner, Serge P
APPLICANT: Vasserot, Allain B
APPLICANT: Vasserot, Allain B
APPLICANT: Pruzan, Ronald A
TITLE OF INVENTION: A Method for Purifying Telomerase
FILE REFERENCE: PurifiedTelomeraseOllbase
CURRENT APPLICATION NUMBER: US/09/717, 828B
CURRENT FILING DATE: 1997-04-06
PRIOR APPLICATION NUMBER: 09/420,056
PRIOR APPLICATION NUMBER: 08/833,377
PRIOR APPLICATION NUMBER: 08/813,377
PRIOR APPLICATION NUMBER: 08/810,736
PRIOR PILING DATE: 1995-08-04
NUMBER: OF SEQ. ID NOS: 11
SOSTWARE: PATENTIN VET. 2.1 edited
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LOCATION: (1)
OTHER INFORMATION: Biotin 5'-terminal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 1, Application US/09717829A Patent No. 6545133
Sequence 1, Application US/09717828B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Artificial Sequence
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US-09-717-829A-1/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 1
LENGTH: 31
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APPLICANT: Weinrich Scott

APPLICANT: Weinrich Scott

APPLICANT: Weinrich Scott

APPLICANT: Weinrich Scott

APPLICANT: Athinson III, Edward

APPLICANT: Alain

APPLICANT: Vasserot, Alain

TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators and

TITLE OF INVENTION: Unhibitors

TITLE OF INVENTION: Unhibitors

TITLE OF INVENTION: Unhibitors

FILE REPREBRUE: 01/062-12-24

CURRENT FILING DATE: 2002-12-24

FRIOR FILING DATE: 1997-04-04

PRIOR APPLICATION NUMBER: 09/420,056

PRIOR FILING DATE: 1999-10-18

PRIOR FILING DATE: 1999-10-18

PRIOR FILING DATE: 1999-10-18

PRIOR APPLICATION NUMBER: 09/717,828

PRIOR FILING DATE: 1090-11-20

NUMBER OF SEQ ID NOS: 11

SOFTWARE: Patentin version 3.1

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; OTHER INFORMATION: Biotin 3'-terminal ; OTHER INFORMATION: Description of Artificial Sequence: Affinity Agent US-09-717-829A-1
                                                                                                                                                                                                                                                                        GapB
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Patent No. 5583016

GENERAL INFORMATION:
APPLICANT: VILLEPONYEAU, Bryant
APPLICANT: FENG, Juni;
APPLICANT: FUNEW, Walter
APPLICANT: ANDREWS, William H.
TITLE OF INVENTION: HUMAN TELOMERASE
NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend Khourie and Crew
STREET: 379 Lytton Avenue
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                      42 TIGICIAACCCIAACTGAGAAGGGCGTAGGC 72
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20;
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                                                                                                                                                                             Query Match 6.9%; Score 31; DB Best Local Similarity 100.0%; Pred. No. 20; Matches 31; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6.9%; Score 31; DB 100.0%; Pred. No. 20; ive 0; Mismatches
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Patent No. 6787133
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Best Local Similarity 100.0
Matches 31; Conservative
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ORGANISM: Homo sapiens
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015389-000830US
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CLASSIFICATION 435
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
FILING DATE: 27-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-UL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-0008
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-660-678A-22/c
; Sequence 22, Application US/08660678A
; Patent No. 5837857
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFRENCE/DOCKET NUMBER: 01536
TELECOMMUNICATION INFORMATION:
TELEPAK: (415) 576-0200
INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
LENGATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
LENGATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
TENGTH: 30 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 30 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                             TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                  linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          , MOLECULE TYPE: DNA
US-08-660-678A-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear
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                                                                                                                                                                                                                                                                                                                             US-08-482-115B-22
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APPLICANT: Feng, Junia
APPLICANT: Feng, Junia
APPLICANT: Funk, Walter
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Assays for the RNA Component of Human
TITLE OF INVENTION: Telomerase
TITLE OF SEQUENCES: 40
CORRESPONDENCE ADDRESS:
ADDRESSEE: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              6.7%; Score 30; DB 1; Length 30; 100.0%; Pred. No. 23; tive 0; Mismatches 0; Indels
                       COMPUTE: 94301

COMPUTE: PASABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: PatentIn Release #1.0, Version #1.25
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/330,123A
FILING DATE: 27-OCT-1994
CLASSFETCATION: A35
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-OLL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 15389-000810
TELEPHONE: (415) 326-2400
TELEPHONE: (415) 326-2400
TELEFAX: (415) 326-2402
INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
LENGTH: 30 base pairs
TYPE: MOLGAL CACH
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/482,115B
FILING DATE: 07-JUN-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               77 TGCTTTTGCTCCCCGCGCGCTGTTTTCTC 106
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COUNTRY: USA
ZIP. 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 22, Application US/08482115B
Patent No. 5776679
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 6.7
Best Local Similarity 100.
Matches 30; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; MOLECULE TYPE: DNA
US-08-330-123A-22
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US-08-482-115B-22/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY:
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Query Match 6.7%; Score 30; DB 1; Length 30; Best Local Similarity 100.0%; Pred. No. 23; Matches 30; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Villeponteau, Bryant
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junli
APPLICANT: Feng, Junli
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS: ADDRESSE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
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Methods for Detecting the RNA Component of Telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1; Length 30;
23;
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APPLICANT: Avillan, Ariel Athena
APPLICANT: Funk, Walter
APPLICANT: Funk, Walter
APPLICANT: Gredder, Carol
APPLICANT: Warhuenda, Maria Antonia Blasco
APPLICANT: Willeponceau, Bryant
TITLE OF INVENTION: RNA COMPONENT OF TELOMERASE
NUMBER OF SEQUENCES: 45
CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: Two Militia Drive
CITY: Lexington
STATE: MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                          COUNTRY: CALL 1984

ZIP: 9411-3834

COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/770,565
FILING DATE: 20-DEC-1996
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/485,778
FILING DATE: 07-JE-1995
                                                                           LLP
TITLE OF INVENTION: Methods for Detecting th
TITLE OF INVENTION: Telomerase
NUMBER OF SEQUENCES: 26
NUMBER OF SEQUENCES: 26
ADDRESSEE: TOWNSEND and TOWNSEND and CREW
STREET TWO Embarcadero Center, 8th Floor
GITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                        015389-002300US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        137 CCTGCCGCCTTCCACCGTTCATCTAGAGC 166
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30 cerececerrecaceerrearreradade 1
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                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INPORMATION:
NAME: Storella, John R.
REGISTRATION UNDRER: 32,94
REFERENCE/DOCKET NUMBER: 01536
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-576-0200
INPORMATION POR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 100
nes 30; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 30 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 02173
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-08-485-778-18/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-770-565-8
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Matches
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   DB 1; Length 30;
23;
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23;
                                       0; Indels
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MEDIUM TYPE: Floppy disk
COMPUTER: Floppy disk
COMPUTER: PLOPPY disk
COMPUTER: THE PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/770,565
FILING DATE: 20-DEC-1996
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      6.7%; Scc...
100.0%; Pred. No. ...
0; Mismatches
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NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-002300US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-576-0200
                                                                           TTCTC 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          290 CTGCCACCGCGAAGAGTTGGGCTCTGTCAG 319
 6.7%; Score 30; DB
100.0%; Pred. No. 23;
tive 0; Mismatches
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Patent No. 5846723
GENERAL INFORMATION:
APPLICANT: Wu, Fred
APPLICANT: Wu, Fred
APPLICANT: Pruzan, Ronald
APPLICANT: Pruzan, Ronald
APPLICANT: Weinrich, Scott L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 100.0
Matches 30; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 30 base pairs
                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: San Francisco
STATE: California
COUNTRY: USA
                                                                           77 TGCTTTTGCTCC
 Query Match
Best Local Similarity
Matches 30; Conserval
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US-08-770-565-8/c
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                                                                                                                                                                                                                          DB 1; Length 30;
23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Andrews, William H.
APPLICANT: Andrews, William H.
APPLICANT: Punk, Walter
APPLICANT: Funk, Walter
APPLICANT: Funk, Walter
APPLICANT: Greider, Carol
APPLICANT: Marhuenda, Maria A. B.
APPLICANT: Marhuenda, Maria A. B.
APPLICANT: Willeponteau, Bryant
TITLE OF INVENTION: RNA Component of Telomerase
NUMBER OF SEQUENCES: 47
CORRESPONDENCE. ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: Two Militia Drive
CITY: Lexington
STATE: Mariallon
                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: TBM PC Compatible
OPERATUR SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/520,550A
FILING DATE: 29-AUG-1995
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PULGATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/387,524
FILING DATE: 13-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-0CT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Granahan, Patricia
REGISTRATION NUMBER: 32,227
REBERENCE/DOCKET NUMBER: CSHL94-05A3B
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                               77 IGCTTTTGCTCCCCGCGCGCTGTTTTCTC 106
                                                                                                                                                                                                                          Query Match 6.7%; Score 30; DB Best Local Similarity 100.0%; Pred. No. 23; Matches 30; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-520-550A-18/c
; Sequence 18, Application US/08520550A
23:
                     SEQUENCE CHARACTERISTICS:
I LENGTH: 30 base pairs
I TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA
US-08-472-802C-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 617-861-9540 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 30 base pairs
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     linear
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US
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US-08-520-550A-18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

MEDIUM TYPE: Floppy disk

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

APPLICATION NUMBER: US/08/472,802C

FILING DATE: 07-40L-1994

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/330,123

RILING DATE: 27-6CT-1994

ATTORNEY/AGENT INFORMATION:

NAME: SMITCH, WILLIAM M.

REGISTRATION NUMBER: 30,223

REFERENCE/DOCKET NUMBER: 15389-000820

TELECOMMONICATION INFORMATION:

TELECOMMONICATION INFORMATION:

TELECOMMONICATION INFORMATION:

TELEFAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 23, Application US/08472802C

Sequence 23, Application US/08472802C

GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junii
APPLICANT: Andrews, William H.

TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 44

CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: TWO Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    77 IGCTITIGCICCCCGCGCGCTGIIITICIC 106
                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/387,524
FILING DATE: 13-FEB-1995
PRIOR APPLICATION DATE: US 08/330,123
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION DATA: APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-OCT-1994
ATTORNEY/AGENT: NFORWATION:
NAME: Granahan, Patricia
REGISTRATION NUMBER: 32,227
REFERENCE/DOCKET NUMBER: CSHL94-05A4
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 1NFORMATION:
TELEFRA: 617-661-6240
INPORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30 recririrecrecedededererrirere 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 30 base pairs;
TYPE: nucleic acid;
TYPE: TRADEDNESS: single
TOPOLOGY: linear
US-08-485-778-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 San Francisco
California
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US-08-472-802C-23/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE:
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STATE: CALIFORNIA
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: PatentIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/060,523
FILING DATE: 14-APR-1998
CLASSIFICATION NUMBER: US/09/060,523
FILING DATE: 05-JUN-1996
RRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
APPLICATION NUMBER: US 08/320,123
FILING DATE: 07-JUL-1994
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
APPLICATION NUMBER: 015389-000813US
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION
                                                                                                                                                                                                                        ADDRESSEE: Townsend and Townsend and Crew LL STREET: Two Embarcadero Center, Bighth Floor CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 6.7%; Score 30; DB Best Local Similarity 100.0%; Pred. No. 23; Matches 30; Conservative 0; Mismatches
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                                    Feng, Junli
Funk, Walter
Addrews, William H.
VENTION: Mammalian Telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: VILLEPONTEAU, Bryant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 379 Lytton Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; Sequence 22, Application US/09580517; Patent No. 6320039; GENERAL INFORMATION:
      Villeponteau, Bryant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        77 IGCITIIGCICCCCGCGCGCGCC
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Palo Alto
STATE: California
APPLICANT: Villeponteau,
APPLICANT: Feng, Junil
APPLICANT: Funk, Walter
APPLICANT: Andrews, Will
ITLE OF INVENTION: Mamm
NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 30 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE:
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                                       0; Indels
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Sequence 22, Application US/08998443

Sequence 22, Application US/0898443

Fatent No. 6054704:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junil
APPLICANT: Funk, Walter
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
APPLICANT: Andrews, William H.
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LIP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/NS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/998,443
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AFFLING DATE:
CLASSIFICATION NUMBER: US/08/599,443
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/660,678
FILING DATE: 05-UN-1996
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-UL-1944
ATTORNEY AGENT INFORMATION:
NAME: SCOREIL OADH R.
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
LENGTH: 130 ABB REPAIRS
   Best Local Similarity 100.0%; Pred. No. 23; Matches 30; Conservative 0; Mismatches
                                                                                                   77 TGCTTTTGCTCCCGCGCGCGCTGTTTTCTC 106
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Patent No. 6258535
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
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US-09-060-523-22/c
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Gaps .. 0 FENG, Junli FUNK, Walter AUDREWS, William H. TITLE OF INVENTION: HUMAN TELOMERASE NUMBER OF SEQUENCES: 25 CORRESPONDENCES: ADDRESS: ADDRESSEE: Townsend and Townsend Khourie and Crew 6.7%; Score 30; DB 1; Length 30; 00.0%; Pred. No. 23; Indela

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dilarity 100.0%; Pred. No. 23; Conservative 0; Mismatches
Best Local Similarity
Matches 30; Conserv
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                                                         MEDIUM TYPE: Floppy disk
COMPUTER: 1BM PC Comparible
COMPUTER: STREM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
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APPLICANT: Weinrich, Scott L
APPLICANT: Ackinson III, Edward M
APPLICANT: Vasserot, Allain P
APPLICANT: Pruzan, Ronald A
ITILE OF INVENTION: A Method for Purifying Telomerase
CURRENT APPLICATION NUMBER: US/09/717,828B
CURRENT PILING DATE: 1999-10-18
FRIOR APPLICATION NUMBER: 09/420,056
FRIOR APPLICATION NUMBER: 08/510,736
FRIOR AP
                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 08/330,123
FILING DATE: cUnknown>
ATTORNEY/AGENT INFORMATION:
NAME: SMith, William REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 15389-000810
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                       CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/580,517
FILING DATE: 25-May-2000
CLASSIFICATION: vUnknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30 TGCTTTTGCTCCCCGCGCGCTGTTTTCTC 1
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SEQUENCE DESCRIPTION: SEQ ID NO: 22:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (415) 326-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 4, Application US/09717828B
Patent No. 6517834
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (415) 326-2422
INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 30 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: single
                    COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: misc_feature
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US-09-717-828B-4/c
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6.7%; Score 30; DB 1; Length 30;

Query Match

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     Gaps
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CRGANISM: Artificial Sequence
CRGANISM: PEATURE:
NAME/KEY: misc_feature
LOCATION: (1)
OTHER INFORMATION: Biotin 5'-terminal
CTHER INFORMATION: Bescription of Artificial Sequence: Affinity Agent
US-09-717-8288-5
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     Indels
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                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Weinrich, Scott L
APPLICANT: Ackinson III, Edward M
APPLICANT: Ackinson III, Edward M
APPLICANT: Ackinson III, Edward M
APPLICANT: Uchteteiner, Serge P
APPLICANT: Vasserot, Allain P
APPLICANT: Prizan, Ronald A
ITITE OF INVENTION: A Method for Purifying Telomerase
CURRENT APPLICATION NUMBER: US/09/717,828B
CURRENT FILING DATE: 1090-10-16
PRIOR APPLICATION NUMBER: 09/420,056
PRIOR APPLICATION NUMBER: 08/833,377
PRIOR APPLICATION NUMBER: 08/833,377
PRIOR FILING DATE: 1997-04-04
PRIOR FILING DATE: 1995-08-04
NUMBER OF SEQ ID NOS: 11
SEQ ID NO SEQ ID NOS: 2.1 edited
IEBNGTH: 30
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| FATCHT NOT. 6549131
| APPLICANT: Weinrich, Scott L
| APPLICANT: Atkinson III, Edward M
| APPLICANT: Atkinson III, Edward M
| APPLICANT: Lichtsteiner, Serge P
| APPLICANT: Lichtsteiner, Serge P
| APPLICANT: Pruzan, Ronald A
| FILE REFERENCE: PurificaTrelomerase011base
| CURRENT APPLICATION NUMBER: US/09/717,829A
| CURRENT APPLICATION NUMBER: 09/420,056
| PRIOR PELING DATE: 1999-10-18
| PRIOR APPLICATION NUMBER: 08/833,377
| PRIOR FILING DATE: 1999-04-04
| PRIOR FILING DATE: 1999-04-04
| PRIOR APPLICATION NUMBER: 08/510,736
| PRIOR APPLICATION NUMBER: 08/510,736
| NUMBER OF SEQ ID NOS: 11
| SOFFWARE: Patentin Ver: 2.1 edited
| SEQ ID NO 4
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                                                  167 AAACAAAAATGTCAGCTGCTGGCCCGTTC 196
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                                                                                                       30 AAACAAAAATGTCAGCTGCTGGCCCGTTC 1
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; Sequence 4, Application US/09717829A
; Patent No. 6545133
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Townsend and Crew LLF

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Townsend and
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STRANDEDNESS: sing
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ADDRESSEE:
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                    TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)
OTHER INFORMATION: Biotin S'-terminal
OTHER INFORMATION: Description of Artificial Sequence: Affinity Agent
US-09-717-829A-4
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OTHER INFORMATION: Description of Artificial Sequence: Affinity Agent
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                                                                                                                                                                                                                                    Query Match 6.7%; Score 30; DB 1; Length 30;
Best Local Similarity 100.0%; Pred. No. 23;
Matches 30; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Machinich, Scott L
APPLICANT: Atkinson III, Edward M
APPLICANT: Lichtereiner, Serge P
APPLICANT: Lichtereiner, Serge P
APPLICANT: Lichtereiner, Serge P
APPLICANT: Pruzan, Rohald A
TILLE OF INVENTION: A Method for Purifying Telomerase
TILLE REFERENCE: Purified-lomerase011base
CURRENT APPLICATION NUMBER: US/09/717,829A
CURRENT FILING DATE: 1909-10-18
PRIOR APPLICATION NUMBER: 08/833,377
PRIOR APPLICATION NUMBER: 08/833,377
PRIOR APPLICATION NUMBER: 08/510,736
PRIOR APPLICATION NUMBER: 08/510,736
PRIOR APPLICATION NUMBER: 08/510,736
PRIOR PILING DATE: 1995-08-04
NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin Ver. 2.1 edited
LENGTH: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 22, Application US/09057351
Patent No. 6548298
GENERAL INFORMATION:
APPLICANT: Feng, Junit
APPLICANT: Feng, Junit
APPLICANT: Funk, Walter
APPLICANT: Funk, Walter
APPLICANT: Mammalian H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
                                                                                                                                                                                                                                                                                                                                     167 AAACAAAATGTCAGCTGCTGCCCGTTC 196
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 137 CCTGCCGCCTTCCACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                   30 AAACAAAAATGTCAGCTGCTGGCCCGTTC 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity 100.
Matches 30; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: misc_feature LOCATION: (1)
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US-09-057-351-22/c
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US-09-903-461-2/C
US-09-903-461-2/C
US-09-903-461-2/C
Sequence 2, Application US/09903461
Fatent No. 6602669
Fatent No. 6602669
FABERALI INFORMATION:
APPLICANT: Letsinger, Robert L.
APPLICANT: Garimella, Viswanadham
TITLE OF INVERTION: Method of Detection by Enhancement of Silver Staining
FILE REFERENCE: 00-1086-A
CURRENT APPLICATION NUMBER: US/09/903,461
CURRENT FILING DATE: 2001-07-11
FRIOR APPLICATION NUMBER: 60/217,782
FRIOR APPLICATION NUMBER: 60/217,782
FRIOR APPLICATION NUMBER: 60/217,782
SOFTWARE: MICCOSOft Word 98
SEQ ID NO 2
LENGTH: 30
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                                                                                                         STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
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; OTHER INFORMATION: synthetic oligomer
US-09-903-461-2
                                                        ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Artificial Sequence
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Best Local Similarity 100.8
Matches 30; Conservative
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OTHER INFORMATION: /note= "N = 5' biotinylated guanosine"
                                                                                                                                                                                                       DB 1; Length 30;
23;
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; OTHER INFORMATION: /note= "oligonucleotide anti-P"
US-08-833-377-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATION SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/833,377
FILING DATE: 04-APR-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATE:
APPLICATION NUMBER: US 08/510,736
FILING DATE: 04-AUG-1995
ATTONINY/AGENT INPORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Weinrich, Scott L.
APPLICANT: Atkinson III, Edward M.
APPLICANT: Lichtsteiner, Serge P.
APPLICANT: Vasserot, Alain P.
APPLICANT: Pruzan, Ronald A.
APPLICANT: Kealey, James T.
TITLE OF INVENTION: Purified Telomerase
NUMBER OF SEQUENCES: 15
CORRESPEDE: ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: Two Embarcadero Center, Bighth Floor
CITY: San Francisco
STATE: California
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                                                                                                                                                                                                    6.7%; Score 30; DB 100.0%; Pred. No. 23; Live 0; Mismatches
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Patent No. 5968506
GENERAL INFORMATION:
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REGISTRATION NUMBER: 32,944
REPRENCE/DOCKET NUMBER: 0153;
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
                                    SOFTWARE: PatentIn version 3.1
SEQ ID NO 5
LENGTH: 30
TYPE: DNA
NORGANIEM: HOMO Sapiens
US-10-330-872-5
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Best Local Similarity 100.(
Matches 30; Conservative
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TYPE: nucleic acid
STRANDEDNESS: single
                NUMBER OF SEQ ID NOS: 11
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NAME/KEY: - 1.30
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Sequence 5, Application US/10330872

Patent No. 6787133

GENERAL INFORMATION:
APPLICANT: Genon Corporation
APPLICANT: Atkinson III, Edward
APPLICANT: Lichtsteiner, Serge
APPLICANT: Lichtsteiner, Serge
APPLICANT: Pruzan, Ronald
ITTLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators and
ITTLE OF INVENTION: Unipp Purified Telomerase to Identify Telomerase Activators and
ITTLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators
FILE REFERENCE: 011/006C

CURRENT FILING DATE: 2002-12-24
PRIOR APPLICATION NUMBER: 08/510,736
PRIOR FILING DATE: 1995-08-04
PRIOR FILING DATE: 1997-04-04

PRIOR FILING DATE: 1997-04-04

PRIOR FILING DATE: 1997-04-04

PRIOR PLING DATE: 1997-04-04

PRIOR PLING DATE: 1997-04-04

PRIOR PLING DATE: 1997-04-04
                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Geron Corporation
APPLICANT: Weinrich, Scott
APPLICANT: Weinrich, Scott
APPLICANT: Weinrich, Scott
APPLICANT: Lichtsteiner, Serge
APPLICANT: Lichtsteiner, Serge
APPLICANT: Lichtsteiner, Serge
APPLICANT: Usersor, Alain
APPLICANT: Usersor, Alain
APPLICANT: Usersor, Alain
APPLICANT: Usersor, Main
APPLICANT: Usersor, Main
FILE REFRENCE: 01/006C
CURRENT APPLICATION INWBER: 08/10/330,872
CURRENT FILING DATE: 2002-12-24
PRIOR APPLICATION NUMBER: 08/833,377
PRIOR FILING DATE: 1995-08-04
PRIOR FILING DATE: 1995-10-18
PRIOR APPLICATION NUMBER: 09/420,056
PRIOR FILING DATE: 1999-10-18
PRIOR APPLICATION NUMBER: 09/717,828
PRIOR APPLICATION NUMBER: 09/717,828
PRIOR APPLICATION NUMBER: 09/717,828
PRIOR FILING DATE: 1000-11-20
NUMBER OF SEQ ID NOS: 11
SCOTTANES: PATCHIL VETSION 3.1
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Pred. No. 23;
                                      0; Indels
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100.0%; Pred. No. 2c.,
0; Mismatches
                                                                                137 CCTGCCGCCTTCCACCGTTCTAGAGC 166
              Pred. No. 23;
100.0%; Preu. ...
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                                                                                                             30 CCTGCCGCCTTCCACCGTTCATTCTAGAGC 1
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APPLICATION NUMBER: 09/717,828
FILING DATE: 2000-11-20
                                                                                                                                                                                                                                             Sequence 4, Application US/10330872
Patent No. 6787133
         Best Local Similarity 100.
Matches 30; Conservative
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Best Local Similarity 100.
Matches 30; Conservative
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ORGANISM: Homo sapiens
                                                                                                                                                                                               RESULT 27
US-10-330-872-4/c
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LENGTH: 30
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/note= "N = 5' biotinylated guanosine"
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28;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER KEADABLE FORM:
MEDIUM TYPE: FI-DOPPY disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/833,377
FILING DATE: 04-APR-1997
CLASSIFICATION NUMBER: US 08/510,736
PRIOR APPLICATION NUMBER: US 08/510,736
FILING DATE: 04-AUG-1995
ATTORNEY/AGENT INFORMATION:
NAME: SLOTELLA, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/POCKET NUMBER: 32,944
REFERENCE/POCKET NUMBER: 32,944
REFERENCE/POCKET NUMBER: 015389-001110US
                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor TTY: San Francisco STATE: California
                                                                                                                                                 APPLICANT: Weinrich, Scott L.
APPLICANT: Atkinson III, Edward M.
APPLICANT: Lichtsteiner, Serge P.
APPLICANT: Vasserot, Alain P.
APPLICANT: Realey, James T.
TITLE OF INVENTION: Purified Telomerase NUMBER OF SEQUENCES: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6.4%; Score 29; DB 100.0%; Pred. No. 28; Live 0; Mismatches
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; Sequence 3, Application US/09717828B
; Patent No. 6517834
                                                                                    Sequence 6, Application US/08833377
Patent No. 5968506
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 30 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 100....
The 29, Conservative
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                       USA
                                            RESULT 31
US-08-833-377-6/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 32
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Query Match 6.4%; Score 29; DB 1; Length 30; Best Local Similarity 100.0%; Pred. No. 28; Matches 29; Conservative 0; Mismatches 0; Indels
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100.0%; Pred. No. 28;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: Callians

COUNTRY: USA

ZIP: 9411-3834

ZIP: 9411-3834

ZEP: 9411-3834

COMPUTER READABLE FORM: Flopty disk mEDIUM TYPE: Flopty disk computable Computable Computable Computable SYSTEM: PEC-DOS/MS-DOS SOFTWARE: Batentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/833,377

FILING DATE: 04-AUG-1997

PRIOR APPLICATION DATA: APPLICATION NUMBER: US 08/510,736

FILING DATE: 132,944

REFERENCY DATE: 32,944

REFERENCY DOCKET NUMBER: 015389-001110US

TELECOWMUNICATION INFORMATION: TELEPHONE: (415) 576-0200

TELEPHONE: CHARACTERISTICS: LENGTH: 30 base pairs

TYPE: nucleic acid

STRANDEDNESS: single
                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Weinrich, Scott L.
APPLICANT: Atkinson III, Edward M.
APPLICANT: Lichtsteiner, Serge P.
APPLICANT: Vasserot, Alain P.
APPLICANT: Prizan, Ronald A.
APPLICANT: Releay, James T.
ITLE OF INVENTION: Purified Telomerase
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
ADDRESSEE: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        167 AAACAAAAATGTCAGCTGCTGGCCCGTT 195
                                                                                         43 TGTCTAACCCTAACTGAGAAGGGCGTAGG 71
                                                                                                                   30 TGTCTAACCCTAACTGAGAAGGGCGTAGG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: -
LOCATION: 1..30
OTHER INFORMATION: /note= "Oligo 13"
                                                                                                                                                                                                                                               Sequence 5, Application US/08833377
Patent No. 5968506
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: modified base
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Best Local Similarity 100.0
Matches 29; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: Two Embarcac
CITY: San Francisco
STATE: California
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Gaps

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Indels

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0; Mismatches

29; Conservative

Matches

à d

412 GAGCTGTGGGACGTGCACCCAGGACTCGGC 441

30 GAGCTATGGGACGTGCACCCAGGACTCGGC 1

Sequence 3, Application US/10330872 Patent No. 6787133 GENERAL INFORMATION: APPLICANT: Geron Corporation

US-10-330-872-3/c

Wed Aug 24 14:55:17 2005

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OTHER INFORMATION: Biotin 5'-terminal OTHER INFORMATION: Description of Artificial Sequence: Affinity Agent
         APPLICANT: Wellicul, SCOLL L
APPLICANT: Atkinson III, Edward M
APPLICANT: Lichteteiner, Serge P
APPLICANT: Vesserot, Allain P
TITLE OF INVENTION: A Method for Purifying Telomerase
CURRENT APPLICATION NUMBER: US/09/17,828B
CURRENT APPLICATION NUMBER: 09/420,056
PRIOR PILING DATE: 1999-10-18
PRIOR APPLICATION NUMBER: 08/833,377
PRIOR APPLICATION NUMBER: 08/833,377
PRIOR APPLICATION NUMBER: 08/833,377
PRIOR FILING DATE: 1999-06-04
NUMBER OF SEQ ID NOS: 11
SOFTWARE: PARCHILING DATE: 1995-08-04
NUMBER OF SEQ ID NOS: 11
SOFTWARE: PARCHILING DATE: 2.1 edited
SEQ ID NOS: 11
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Patent No. 6545133

GENERAL INCORMATION:
APPLICANT: Weinrich, Scott L
APPLICANT: Lichtetiener, Serge P
APPLICANT: Pruzan, Ronald A
TITLE OF INVENTION: A Method for Purifying Telomerase
CURRENT APPLICATION NUMBER: US/09/717,829A
CURRENT APPLICATION NUMBER: 09/420,056
PRIOR PILING DATE: 1999-10-18
PRIOR PELLING DATE: 1999-04-04
PRIOR FILING DATE: 1999-04-04
PRIOR FILING DATE: 1999-04-04
PRIOR FILING DATE: 1995-04-04
PRIOR PILING DATE: 1995-04-04
PRIOR PILING DATE: 1995-08-04
SPRIOR APPLICATION NUMBER: 08/510,736
NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin Ver: 2.1 edited
SEQ ID NO 3
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ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
APPLICANT: Weinrich, Scott L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)
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Best Local
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APPLICANT: Meinrich, Scott
APPLICANT: Aktinson III, Edward
APPLICANT: Aktinson III, Edward
APPLICANT: Lichtsteiner, Serge
APPLICANT: Lichtsteiner, Serge
APPLICANT: Pruzan, Ronald
TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators and
TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators and
TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators and
TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators and
TITLE OF INVENTION: Using Purified Date: 2002-12-24
PRIOR APPLICATION NUMBER: 08/510,736
PRIOR FILING DATE: 1997-04-04
PRIOR PELING DATE: 1997-04-04
PRIOR PELING DATE: 1997-04-04
PRIOR FILING DATE: 1997-04-04
PRIOR FILING DATE: 2000-11-20
NUMBER OF SEQ ID NOS: 11
SOFTWARRE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Johns Hopkins University
APPLICANT: Johns Hopkins University
APPLICANT: Gender, Carol W.
APPLICANT: Le, Siyuan
ITILE OF INVENTION: TELOMERASE-ASSOCIATED PROTEINS
FILE REFERENCE: 07265/157001
CURRENT APPLICATION NUMBER: US/09/286,959B
CURRENT PLING DATE: 1999-04-06
PRIOR APPLICATION NUMBER: 60/080,783
PRIOR PLING DATE: 1998-04-06
NUMBER OF SEQ ID NOS: 24
SOFTWARE: FASCISO FOR Windows Version 4.0
SEQ ID NO 3
ILENGTH 28
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 6.3%; Score 28.4; DB 1;
Best Local Similarity 96.7%; Pred. No. 31;
Matches 29; Conservative 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      412 GAGCTGTGGGACGTGCACCCAGGACTCGGC 441
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; OTHER INFORMATION: Primer US-09-286-959B-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-330-872-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
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OTHER INFORMATION: Biotin 5'-terminal
OTHER INFORMATION: Description of Artificial Sequence: Affinity Agent
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                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1; Length 30;
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Pred. No. 31;
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6.3%;

Query Match Best Local Similarity

US-09-717-829A-3

NAME/KEY: misc_feature

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APPLICANT: Kim, Nam Woo
APPLICANT: Kim, Nam Woo
APPLICANT: Wu, Fred
APPLICANT: Weinrich, Scott L.
APPLICANT: Pruzan, Ronald
APPLICANT: Weinrich, Scott L.
TITLE OF INVENTION: Methods for Detecting the RNA Component of
TITLE OF INVENTION: Telomerase
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: TOWNSEND and TOWNSEND and CREW LLP
STREET: Two Embarcadero Center, 8th Floor
CITY: San Francisco
STRATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-974-180-32/c

Sequence 32, Application US/08974180

Patent No. 6025194

Patent No. 6025194

TITLE OF INVENTION: Methods for Modulating and Identifying TITLE OF INVENTION: Callular Senescence

TITLE OF INVENTION: Callular Senescence

NUMBER OF SEQUENCES: 36

CORRESPONDENCE ADDRESS:

ADDRESSEE: Geron Corporation

STREET: 230 Constitution Drive

CITY: Menlo Park

STATE: California

COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1; Length 27;
35;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: Eloppy disk
COMPUTER: Eloppy disk
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
COMPUTER: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/770,565
FILING DATE: 20-DEC-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INPOMMATION:
NAME: Storella, John R:
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 32,944
REFERENCE/OFFET NUMBER: 32,944
RELEPRAK: 415-576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 6.0%; Score 27; DB Best Local Similarity 100.0%; Pred. No. 35; Matches 27; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   144 CCTTCCACCGTTCATTCTAGAGCAAAC 170
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                                                                           Sequence 26, Application US/08770565; Patent No. 5846723; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: 26: SEQUENCE CHARACTERISTICS: LENGTH: 27 base pairs TYPE: nucleic acid STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 94025
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TOPOLOGY: linear; MOLECULE TYPE: DNA US-08-770-565-26
                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
                                RESULT 37
US-08-770-565-26/c
                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
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/note= "N = 5' biotinylated guanosine"
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                                   0; Indels
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SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/833,377

FILING DATE: 04-APR-1997

CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/510,736
FILING DATE: 04-AUG-1995
ATTORNEY/AGENT INFORMATION:
NAME: SLOCABILA, UDAN R.
REGISTRATION NUMBER: 015389-001110US
TELEPHONE: (415) 576-0300
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 4:
SEQUIBNE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                  APPLICANT: Advances
APPLICANT: Advances
APPLICANT: Atkinson III, Edward M.
APPLICANT: Lichtsteiner, Serge P.
APPLICANT: Lichtsteiner, Serge P.
APPLICANT: Lichtsteiner, Serge P.
APPLICANT: Forese, Alain P.
APPLICANT: Fealey, James T.
TITLE OF INVENTION: Purified Telomerase
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
COUNTRY: USA
           Pred. No. 31;
100.0%; Pred. ...
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LOCATION: 1..30
OTHER INFORMATION: /note= "Oligo 5"
                                                                             17 GCCTGGGAGGGGTGGTGGCCATTTTTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                            Sequence 4, Application US/08833377
Patent No. 5968506
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: modified_base LOCATION: 1
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Best Local Similarity 96.6
Matches 28; Conservative
         Best Local Similarity 100.
Matches 28; Conservative
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TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: DNA
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Gaps

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Gaps

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Sequence 40, Application US/08482115B
Fatent No. 5776679
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junii
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Telomerase
TITLE OF INVENTION: Telomerase
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                        5.9%; Score 26.4; DB 1; Length 28; 96.4%; Pred. No. 41; Live 0; Mismatches 1; Indels
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COUNTRY: USA

ZIP: 94111-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/482,115B
FILING DATE: 07-JUN-1995
CLASSIFICATION NUMBER: US/08/482,115B
FILING DATE: 07-JUL-1994
PRICK APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
PRICK APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
ATTORNEY/AGENT INFORMATION:
NAME: STORMATION: NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000830US
TELECOMMUNICATION INFORMATION:
              015389-000830US
                                                                                                                                                                                                                                                                                                                                                                                                                      17 GCCTGGGAGGGGTGGTGGCCATTTTTG 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 GCCTGGGAGGGTGGTGCTATTTTG 28
                                                      TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 31:
SEQUENCE CHARACTERISTICS:
LENGTH: 28 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 40:
SEQUENCE CHARACTERISTICS:
LENGTH: 28 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                  TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
            REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 96.4'
Matches 27; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              San Francisco
: California
                                                                                                                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: DNA
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MOLECULE TYPE:
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Fatent No. 5776679
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Feng, Juni
APPLICANT: Peng, Juni
TITLE OF INVENTION: Assays for the RNA Component of Human
TITLE OF INVENTION: Telomerase
NUMBER OF SEQUENCES:
OORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6.0%; Score 27; DB 1; Length 27;
100.0%; Pred. No. 35;
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ZIP: 94111-3834

COUNTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
SOFTWARE: PATENTIN DATA:
APPLICATION NUMBER: US/08/482,115B
FILING DATE: 07-UNH-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/22,102
FILING DATE: 07-ULL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: Z7-CCT-1994
PRIOR APPLICATION NUMBER: 32,944
FRGISTRATION NUMBER: 32,944
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: -
LOCATION: 1..27
OTHER INFORMATION: /note= "primer hTR445 comp"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Pred. ....
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                                               SOFTWARE: Patentin Release #1.0, Ver CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/974,180
FILING DATE: 19-NOV-1997
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Kaster, Kevin R.
REGISTRATION NUMBER: 206
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 473-779
TELEFAX: (650) 473-8654
INFORMATION FOR SEQ ID NO: 32:
SEQUENCE CHARACTERISTICS:
LENGTH: 27 base pairs
TYPE: nucleic acid
STRANDENNES: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              27 TGCACCCAGGACTCGGCTCACACATGC 1
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Matches 27; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-482-115B-31
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Sequence 23, Application US/08482115B
Sequence 23, Application US/08482115B
Sexeral INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Punk, Walter
APPLICANT: Peng, Villiam H.
ITILE OF INVENTION: Telomerase
NUMBER OF SEQUENCES: 40
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1; Length 26;
40;
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                                                                                                                                                                                                                           COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

SOFTWARE: PAPLICATION DATA:

APPLICATION NUMBER: US/08/330,123A

FLING DATE: 27-OCT-1994

CLASSIFICATION NUMBER: US 08/272,102

FLING DATE: 07-DUL-1994

ATTORNEY AGENT INPORMATION:

NAME: Smith, William M

REGISTRATION NUMBER: 15389-000810

TELERDANCE/DOCKET NUMBER: 15389-000810

TELERDANCE/OCKET NUMBER: 15389-000810
APPLICANT: ANDREWS, William H.
TITLE OF INVENTION: HUMAN TELOMERASE
COMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend Khourie and Crew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 5.8%; Score 26; DB Best Local Similarity 100.0%; Pred. No. 40; Matches 26; Conservative 0; Mismatches
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                                                                                                                         STREET: 379 Lytton Avenue CITY: Palo Alto STATE: California COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 26 base par
TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       , MOLECULE TYPE: DNA
US-08-330-123A-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    JS-08-482-115B-23/c
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                                   1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PATEM: PC-DOS/MS-DOS
SOFTWARE: PATEMIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/472,802C
FILING DATE: 07-UNN-1995
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                       APPLICANT: VIleponteau, Bryant
APPLICANT: Feng, Junii
APPLICANT: Feng, Junii
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 44
CORRESPONDENCE ADDRESS: AND ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION 514
PELORASIFICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-UUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-0CT-1994
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
RESPERENCE/DOCKET NUMBER: 15389-000820
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 38:
ERRIGHT: 28 base pairs
          Pred. No. 41;
9; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                      17 GCCTGGGAGGGGTGGTGGCCATTTTTG 44
                                                                                                                    1 GCCUGGGAGGGUGGUGGCUAUUUUUG 28
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Sequence 23, Application US/08330123A
Petent No. 5583016
Petent No. 5583016
PETENT NIPORMATION:
APPLICANT: FENG, Juni;
APPLICANT: PENG, Juni;
APPLICANT: PENG, Juni;
APPLICANT: PENG, Walter
                                                                                                                                                                                                                                                              ; Sequence 38, Application US/08472802C; Patent No. 5958680; GENERAL INFORMATION:
          64.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches 27; Conservative
          Best Local Similarity 64.3
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: Two bulled CITY: San Francisco CTATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
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Best Local Similarity
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US-08-472-802C-38
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                                                                                                                                                                                                                                                                                          0; Indels
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Pred. No. 40;
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COUNTRY: USA
ZIP: 9411-3834

CZIP: 9411-3834

CZIP: 9411-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Parentin Release #1.0, Version #1.30

SOFTWARE: Parentin Release #1.0, Version #1.30

CURENT APPLICATION DATA:
APPLICATION NUMBER: US/08/660,678A

FILING DATE: 05-UJN-1996

CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123

FILING DATE: 27-OCT-1994

PRIOR APPLICATION DATA:
APPLICATION NUMBER: 32,944

REFERENCE STORELLA, JOHN R.
REFILING DATE: 07-UJL-1994

ATTORNEY/AGENT INFORMATION:
NUMBER: STORELLA, JOHN R.
REFILING DATE: 07-UJL-1994

ATTORNEY/AGENT INFORMATION:
TELEPONE: (415) 576-0300

TELEFAX: (415) 576-0300

TELEFAX: (415) 576-0300

INFORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
LENGTH: C2 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Mammalian relomerase
NUMBER OF SEQUENCES: 30
CARRESPONDENCE ADDRESS:
ADDRESSEB: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5.8%; Scc. 100.0%; Pred. No. 20.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                         45 TCTAACCCTAACTGAGAAGGGCGTAG 70
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US-08-660-678A-23/c
; Sequenc 23, Application US/08660678A
; Patent No. 5837857
                      INFORMATION FOR SEQ ID NO: 2
SEQUENCE CRARACTERISTICS:
LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDENESS: single
TOPOLLGY: linear
MOLECULE TYPE: DNA
  (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 100.0
Matches 26; Conservative
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  TELEFAX:
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APPLICANT: Feng, Junli
APPLICANT: Feng, Junli
APPLICANT: Funk, Walter
APPLICANT: Andrews, William H.
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Assays for the RNA Component of Human
TITLE OF INVENTION: Telomerase
NUMBER OF SEQUENCES: 40
CORRESPONDENCE ADDRESS: 40
CORRESPONDENCE ADDRESS: ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 26; DB 1; Length 26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CAPPUTER FRADABLE FORM:
MEDIUM TYPE: Floppy disk
CAMPUTER ERADABLE FORM:
COMPUTER: END FO Compatible
OPERATING SYSTEM: FC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
FILING DATE: 07-JUL-1994
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-CCT-1994
ATTORNEY/AGENT INFORMATION:
NAME: SCOTELLA, JOHN R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 32,944
REFERENCE/COMMUTICATION INFORMATION:
TELECOMMUTICATION INFORMATION:
               CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNDRER:
FILING DATE: 07-011-1994
PRIOR APPLICATION UNDRER: US 08/272,102
PRIOR APPLICATION UNDRER: US 08/330,123
PILING DATE: 27-007-1994
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000830US
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDENNES: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 5.8%; Score 26; DB Best Local Similarity 100.0%; Pred. No. 40; Matches 26; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CTTCCACCGTTCATTCTAGAGCAAAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 29, Application US/08482115B Patent No. 5776679 GENERAL INFORMATION:
07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: USA
ZIP: 94111-3834
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-482-115B-23
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US-08-482-115B-29
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PRIOR APPLICATION DATA:
       San Francisco
                             California
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US-08-710-249-25
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                                                                                                                                                                                                                        APPLICANT: Wu, Fred
APPLICANT: Kealey, James T.
APPLICANT: Pruzan, Ronald
APPLICANT: Pruzan, Ronald
APPLICANT: Pruzan, Ronald
APPLICANT: Pruzan, Ronald
APPLICANT: Weinran, Scott L.
TITLE OF INVENTION: Telomerase
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: TOWNSEND and TOWNSEND and CREW LLP
STREET: Two Embarcadero Center, 8th Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 25, Application US/08710249

Patent No. 5858777

GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Andrews, William H.
APPLICANT: Andrews, Wethods and Telomerase Activity
NUMBER OF SEQUENCES: 26

CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CUDNIKY: U.S.A.

ZIP: 94111-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/770,565
FILING DATE: 20-DEC-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: SCOTELLA, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET UNMBER: 015389-002300US
TELEFHONE: 415-576-0200
TELEFAX: 415-576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      E: Townsend and Townsend and Crew LLP
Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             60 GAAGGCCTAGGCGCCCTGCTTTTGC 85
26 CTTCCACCGTTCATTCTAGAGCAAAC 1
                                                                                                                                        Sequence 25, Application US/08770565 Patent No. 5846723 GENERAL INFORMATION: GENERAL INFORMATION: APPLICANT: Kim, Nam Woo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5.5°, 100.08°, 11°, 0°,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 415-576-0300
INFORMATION FOR SEQ ID NO: 5
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
Matches 26; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: DNA
                                                                                            RESULT 46
US-08-770-565-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-08-710-249-25
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                                                                                                                                                                                                                                                                                                                                                                             DB 1; Length 26; 40;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/472,802C FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Vileponteau, Bryant
APPLICANT: Feng, Junii
APPLICANT: Feng, Junii
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 44
CORRESPONDENCE ADDRESS: ANDRESSE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PULLASSIFILION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 15389-000820
TELECOMMUNICATION: NECRMATION:
                                                                                                                                                                                                                                                                                                                                                                             Query Match 5.8%; Score 26; DB Best Local Similarity 100.0%; Pred. No. 40; Matches 26; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  145 CTTCCACGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        145 CTTCCACCGTTCATTCTAGAGCAAAC 170
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: Two Embarcadero Center, Bi
CITY: San Prancisco
STATE: California
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC CO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-472-802C-24/c
; Sequence 24, Application US/08472802C
; Patent No. 5958680
; GENERAL INFORMATION:
          TELECOMMUNICATION INFORMATION:
                                                                          TELEFAX: 617-861-9540 INFORMATION FOR SEQ ID NO: 19:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (415) 576-0200
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INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26; Conservative
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                                                                                                                                                                                                                                                                                 ; TOPOLOGY: linear
US-08-485-778-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 26; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-08-472-802C-24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Villeponteau, Bryant
TITLE OF INVENTION: RNA COMPONENT OF TELOMERASE
NUMBER OF SEQUENCES: 45
CORRESPONDENCES: 45
ADDRESSEB: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: Two Militia Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: RIOPRY
COMPUTER: IRANGE READABLE FORM:
MEDIUM TYPE: RIOPRY
COMPUTER: IBM PC compatible
COMPUTER: BEATHING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
SUGNERNY APPLICATION DATA:
APPLICATION NUMBER: US/08/485,778
FILING DATE: 07-15-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/387,524
FILING DATE: 13-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 13-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION DATA:
ATTORNEY/AGENT INFORMATION:
APPLICATION NUMBER: US 08/583,808
FILING DATE: 05-JAN-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/003,492
FILING DATE: 08-SEP-1995
ATORNEY/AGENT INFORMATION:
NAME: Stochl3, John R.
REGISTRATION NUMBER: 31,944
REGISTRATION NUMBER: 31,944
REFERENCE/DOCKET NUMBER: 015389-001220US
TELECOMMUNICATION INFORMATION:
TELEPAN: (415) 576-0300
TELEPAN: (415) 576-0300
INFORMATION FOR SEQ ID NO: 26: SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
FENCHH: 26 base pairs
TYPE: nucleic acid
STRANDEDMES: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Greider, Carol
Marhuenda, Maria Antonia Blasco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Granahan, Patricia
REGISTRATION NUMBER: 32,227
REFERENCE/DOCKET NUMBER: CSHL94-05A4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             145 CTTCCACCGTTCATTCTAGAGCAAAC 170
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Avilion, Ariel Athena
Feng, Junli
Funk, Walter
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Patent No. 5876979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 26; Conserv
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US-08-485-778-19/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-710-249-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
APPLICANT:
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US-08-974-180-33

Sequence 33, Application US/08974180

Patent No. 602519.

Patent No. 602519.

TILLE OF INVENTION: Methods for Modulating and Identifying TITLE OF INVENTION: Callular Senescence

TITLE OF INVENTION: Callular Senescence

TITLE OF INVENTION: Callular Senescence

NUMBER OF SEQUENCES: 36

CORRESPONDERCE: 36

CORRESPONDERCE: Geron Corporation

STREET: 230 Constitution Drive

CITY: Menlo Park

STATE: Galifornia

COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5.8%; Score 26; DB 1; Length 26; ilarity 100.0%; Pred. No. 40; Conservative 0; Mismatches 0; Indels
TITLE OF INVENTION: RNA Component of Telomerase NUMBER OF SEQUENCES: 47
CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C. STREET: Two Militia Drive
CITY: Lexington
STATE: MA
                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: PatentIn Release #1.0, Version #1.30
SOCTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/520,550A
FILING DATE: 29-AUG-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/387,524
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/320,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/372,102
FILING DATE: US 08/372,102
ATTOMENY/AGENT INFORMATION:
ANAMER OF THE PRIOR DATE: US 08/272,102
ATTOMENY/AGENT INFORMATION:
ANAMER OF THE PRIOR DATE: US 08/272,102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIF: 94025
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Granahan, Patricia
REGISTRATION NUMBER: 32,227
REFERENCE DOCKET NUMBER: CSHL94-05A3B
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-861-6240
INFORMATION FOR SEQ ID NO: 19:
INFORMATION FOR SEQ ID NO: 19:
LENGTH: 26 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 26; Conserv
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US-08-520-550A-19
                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         g
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: ISM PC COMPATIBLE

COMPUTER: ISM PC COMPATIBLE

COMPOTER: ISM PC COMPATIBLE

COMPOTER: ISM PC COMPATIBLE

COMPOTER: ISM PC COMPATIBLE

SOFTWARE: Patentin Release #1.0, Version #1.30

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/472,802C

FILING DATE: 07-JUN-1994

PRIOR APPLICATION NUMBER: US 08/22,102

FILING DATE: 07-JUL-1994

ATTORNEY/AGENT INPORMATION:

NAME: SMATCH, William M.

REGISTRATION NUMBER: 15389-000820

TELEGORMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:
                                                                                                                                       Sequence 30. Application US/08472802C
Patent No. 5958680
GRNERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Juni;
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 44
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                               STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          45 TCTAACCCTAACTGAGAGGGCGTAG 70
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26 CTTCCACGGTTCATTCTAGAGCAAAC 1
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Avilion, Ariel A.
Feng, Junli
Funk, Walter
Greider, Carol
Marhuenda, Maria A. B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19, Application US/08520550A Patent No. 6013468 GENERAL INFORMATION:
APPLICANT: Andrews, William H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Villeponteau, Bryant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (415) 576-0200
INFORMATION FOR SEQ ID NO: 30:
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 100.0
Marches 26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 52
US-08-520-550A-19/c
                                                                                                  RESULT 51
US-08-472-802C-30
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APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT: Lingner, Joachim
APPLICANT: Nakamura, Toru
APPLICANT: Nakamura, Toru
APPLICANT: Chapman, Karen B.
APPLICANT: Harley, Calvin B.
APPLICANT: Harley, Calvin B.
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/974,549A
FILING DATE: 19-NOV-1997
CLASSIFICATION DATA:
APPLICATION DATA: 08/724,643
FILING DATE: 01-OCT-1996
PILING DATE: 18-APR-1997
PILING DATE: 18-APR-1997
PILING DATE: 18-APR-1997
PILING DATE: 18-APR-1997
PILING DATE: 25-APR-1997
PILING DATE: 25-APR-1997
PILING DATE: 25-APR-1997
PILING DATE: 25-APR-1997
PILING DATE: 06-MAY-1997
PILING DATE: US-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                              Query Match
5.8%; Score 26; DB
Best Local Similarity 100.0%; Pred. No. 40;
Matches 26; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                   145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                      26 CTTCCACGTTCATTCTAGAGCAAAC 1
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AUG-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/911,312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 597, Application US/08974549A Patent No. 6166178 GENERAL INFORMATION:
             ; TELEFAX: (415) 576-0300
; INFORMATION FOR SEQ ID NO: 23:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 26 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: DNA
US-08-998-443-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 55
US-08-974-549A-597
                                                                                                                                                                                                                                                                              Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 23, Application US/08998443
; Sequence 23, Application US/08998443
; Patent No. 6054575
; GENERAL INFORMATION:
; APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junil
; APPLICANT: Funk, Walter
; APPLICANT: Andrews, William H.
: TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
ZIP: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/08/998,443
FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; NAME/KEY: -
; LOCATION: 1..26
; CTHER INFORMATION: /note= "primer hTR S328"
US-08-974-180-33
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APPLICATION NUMBER: US/08/660,678
FILING DATE: 05-UN-1996
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-CCT-1994
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-UL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERNEK/DOCKET NUMBER: 32,944
REFERNEK/DOCKET NUMBER: 015389-0008
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
APPLICATION NUMBER: US/08/974,180
                                                                                          ALTORUMY TREAST TREASTACEN
NAME: Kagter, Kevin R.
REGISTRATION NUMBER: 32,704
REFRENCE/DOCKET NUMBER: 206
TELECOMMUNICATION INFORMATION:
TELEPAX: (650) 473-7779
TELEPAX: (650) 473-8654
INFORMATION FOR SEQ ID NO: 33:
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                           FILING DATE: 19-NOV-1997
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-998-443-23/c
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ch 5.8%; Score 26; DB 1; Length 26; 1 Similarity 100.0%; Pred. No. 40; 26; Conservative 0; Mismatches 0; Indels
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                                                                                                                  PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 08/854,050
FILING DATE: 09-MAY-1997
PRIOR APPLICATION NUMBER: US 08/911,312
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/911,312
FILING DATE: 14-A0C-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-A0C-1997
PRIOR APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-A0C-1997
PRIOR APPLICATION NUMBER: WO PCT/US97/17618
FILING DATE: 01-OCT-1997
PRIOR APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-OCT-1997
ATTORNEY AGENT INFORMATION:
APPLICATION NUMBER: 36,429
REGERENCE/DOCKET NUMBER: 36,429
REGERENCE/DOCKET NUMBER: 36,429
RETERENCE/DOCKET NUMBER: 36,429
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; LOCATION: 1..26
; OTHER INFORMATION: /note= "R3c primer"
US-08-974-549A-598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junli
APPLICANT: Fenk, Walter
APPLICANT: Funk, Walter
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         145 CTTCCACCGTTCATTCTAGAGCAAAC 170
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Patent No. 6258535
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 26 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              JS-09-060-523-23/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Cech, Thomas R.
APPLICANT: Linguer, Joachim
APPLICANT: Nakamura, Toru
APPLICANT: Chapman, Karen B.
APPLICANT: Morin, Gregg B.
APPLICANT: Harley, Calvin B.
APPLICANT: Andrews, William H.
TITLE OF INFORTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: San Francisco
CITY: California
CONTRY: USA
ZIP: 94111-3834
ZIP: 94111-3834
ZOMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
COMPOTER: IBM PC COMPATIBLE
SOFTWARE: PAPLICATION DATA:
APPLICATION NUMBER: US/08/974,549A
FILING DATE: 01-0CT-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/844,419
FILING DATE: 19-0CT-1996
PRIOR APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-0AP-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/846,017
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        B: Townsend and Townsend and Crew LLP
Two Embarcadero Center, Eighth Floor
PRIOR APPLICATION DATA:

APPLICATION NUMBER: WO PCT/US97/17618
FILING DATE: 01-0CT-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-0CT-1997
ATORNEY/AGENT INFORMATION:
NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002610US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 597:
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: -
LOCATION: 1..26
OTHER INFORMATION: /note= "F3b primer"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             45 TCTAACCCTAACTGAGAGGGCGTAG 70
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2008-974-549A-59B/c
; Sequence 598, Application US/08974549A
; Patent No. 6166178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend a
STREET: Two Embarcade
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-974-549A-597
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US-09-220-157A-26/C

Sequence 26, Application US/09220157A

Sequence 26, Application US/09220157A

Patent No. 6300110

APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junli
APPLICANT: Addrews, William H.

APPLICANT: Addrews, William H.

TITLE OF INVENTION: Methods and Reagents for Regulating
TITLE OF INVENTION: Telomere Length and Telomerase Activity
NUMBER OF SEQUENCES: 26

CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    5.8%; Score 26; DB 1; Length 26; 100.0%; Pred. No. 40; trive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/220,157A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco
                         FILING DATE: 08-SEP-1995
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001220US
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDEDNES: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          015389-001220US
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CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/710,249
FILING DATE: 13-SEP-1996
APPLICATION NUMBER: US 08/583,808
FILING DATE: 05-JAN-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/003,492
FILING DATE: 08-SEP-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        45 TCTAACCCTAACTGAGAAGGGCGTAG 70
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       US 60/003,492
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 94111.3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
CMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ALTOKNET, AGENT IN CORFACTORY
NAME: SCORELIA, JOHN R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 01531
TELEPHONE: (415) 576-0200
INFORMATION FOR SEQ ID NO: 26:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity 100.0
Matches 26; Conservative
       APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  California
                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: DNA
US-09-220-157A-25
                                                                                                                                                                                                                                                                                                                                                       linear
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                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junii
APPLICANT: Andrews, William H.
APPLICANT: Adams, Robert R.
ITILE OF INVENTION: Methods and Reagents for Regulating
TITLE OF INVENTION: Telomere Length and Telomerase Activity
NUMBER OF SEQUENCES: 26
CORRESPONDENCES: 26
CORRESPONDENCES: ADDRESS: ADDRESSE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
COUNTRY: USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ISP PC compatible
CURENT APPLICATION DATA:
APPLICATION NUMBER: US/09/220,157A
FLIING DATE:
APPLICATION NUMBER: US/08/710,249
FILING DATE: 1.5EP.1996
FILING DATE: 05-JAN-1996
SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                            015389-000813US
                                                                     CLASSIETCATION. 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/660,678
FILING DATE: 05-UNU-1996
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-UL-1994
PRIOR APPLICATION NUMBER: US 08/272,102
PRIOR APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-UL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 22,944
REFERENCE/DOCKET NUMBER: 015389-0006
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFAX: (415) 576-0300
TELERAX: (415) 576-0300
TELEFAX: LENGTH: 26 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26 CTTCCACCGTTCATTCTAGAGCAAAC 1
                                         UMBER: US/09/060,523
14-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 25, Application US/09220157A Patent No. 6300110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
                                              APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; MOLECULE TYPE: DNA
US-09-060-523-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 58
US-09-220-157A-25
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CTTCCACCGTTCATTCTAGAGCAAAC 1
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GENERAL INFORMATION:
                                            <Unknown>
FILING DATE: 25-May-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 26 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-912-951-311
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Sequence 23, Application US/09580517
Sequence 23, Application US/09580517
Setting Time Of 1000 Nation of
                                                                                                                                                                                                                                5.8%; Score 26; DB 1; Length 26;
100.0%; Pred. No. 40;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 26;
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COMPUTER: ISM PC Compatible
CERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 60
US-09-286-9598-4/C
Sequence 4, Application US/09286959B
Patent No. 6300131
GENERAL INFORMATION:
APPLICANT: Johns Hopkins University
APPLICANT: Johns Hopkins University
APPLICANT: Le, Siyuan
TILE OF INVENTION: TELOMERASE-ASSOCIATED PROTEINS
FILE REFERENCE: 07265/157001
CURRENT APPLICATION NUMBER: US/09/286,959B
CURRENT FILING DATE: 1999-04-06
PRIOR APPLICATION NUMBER: 60/080,783
PRIOR FILING DATE: 1998-04-06
NUMBER OF SEQ ID NOS: 24
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4
LENGTH: 26
TURDET FILE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 5.8%; Score 26; DB Best Local Similarity 100.0%; Pred. No. 40; Matches 26; Conservative 0; Mismatches
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APPLICATION NUMBER: US/09/580,517
                                                                                                                                                                                                                                                                                                                                                                                             145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      145 CTTCCACCGTTCATTCTAGAGCAAAC 170
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 94301
COMPUTER READABLE FORM:
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STATE: California
                                                                                                                                                                                                                                                                        Best Local Similarity 100.
Matches 26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Primer US-09-286-959B-4
TYPE: nucleic acid
STRANDEDNESS: Bingl
TOPOLOGY: linear
MOLECULE TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-580-517-23/c
                                                                                                                            ; MOLECULE TYP!
US-09-220-157A-26
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APPLICANT: Lingner, occ...
APPLICANT: Chapman, Karen B.
APPLICANT: Chapman, Karen B.
APPLICANT: Chapman, Karen B.
APPLICANT: Chapman, Karen B.
APPLICANT: Andrews, William H.
ITLE OF INVENTION: HUMAN TELOMERASE CATALYTIC SUBUNIT: DIAGNOSTIC AND TITLE OF INVENTION: HUMAN TELOMERASE CATALYTIC SUBUNIT: DIAGNOSTIC AND TITLE OF INVENTION: THERAPEUTIC METHODS
NUMBER OF SEQUENCES: 335
CORRESPONDENCE ADDRESS:
ADDRESSE: Two sends and Townsend and Crew LLP
STREET: Two Embarcadero Center, 8th Floor
CITY: San Francisco
STATE: California
COUNTRY: United States of America
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MEDIUM TYPE: Floppy disk
COMPUTER: INP Compatible
COMPUTER: INP PC compatible
SOFTWARE: Patentin Release #1.0, Version #1.30
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US 08/854,050
FILING DATE: 09-MAY-1997
CLASSIFICATION NUMBER: US 08/851,843
PRIOR APPLICATION NUMBER: US 08/851,843
PLING DATE: 06-MAY-1997
CLASSIFICATION NUMBER: US 08/851,843
PRIOR APPLICATION NUMBER: US 08/851,843
PRIOR APPLICATION NUMBER: US 08/851,845
FILING DATE: 25-APR-1997
FILING DATE: 25-APR-1997
PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/330,123
FILING DATE: -CURhown-
ATTORNEY/AGRY INFORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 15389-000810
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 326-2420
INFORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
40;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 5.8%; Score 26; DB Best Local Similarity 100.0%; Pred. No. 40; Matches 26; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                         ; TOPOLOGY: linear
; MOLECULE TYPE: DNA
; SEQUENCE DESCRIPTION: SEQ ID NO: 23:
US-09-580-517-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        145 CTTCCACCGTTCATTCTAGAGCAAAC 170
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5.8%; Score 26; DB 1; Length 26; 100.0%; Pred. No. 40; tive 0; Mismatches 0; Indels
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APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
CLASSIFICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
PRIOR APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US 08/310,123
FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California COUNTRY: USA
                                       PRIOR APPLICATION DATA:

PRIOR APPLICATION HUMBER: US 08/846,017
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/724,643
APPLICATION NUMBER: US 08/724,643
APPLICATION INTORATION:
APPLICATION NUMBER: US 08/724,643
ATORNEY/AGENT INFORMATION:
NAMME: APPLICATION NUMBER: 36,429
REGISTRATION NUMBER: 36,429
REPERENCE/DOCKET NUMBER: 015389-002600US
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 312:
SEQUENCE CHARACTERISTICS:
LENGTH : 26 DASE PARIST
MUMBER: MUMBE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 23, Application US/09057351
Patent No. 6548298
GENERAL INFORMATION:
PAPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junii
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES:
CRESPONDENCE ADDRESS: 42
ADDRESSEE: Townsend and Crew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                145 CTTCCACCGTTCATTCTAGAGCAAAC 170
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity 100.
Matches 26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-057-351-23/c
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GENERAL INCORATION:
APPLICANT: Cech, Thomas R.
APPLICANT: Linguer, Joachim
APPLICANT: Chapman, Karen B.
APPLICANT: Chapman, Karen B.
APPLICANT: Harley, Calvin
APPLICANT: Harley, Calvin
APPLICANT: Harley, Calvin
APPLICANT: HUMAN TELOMERASE CATALYTIC SUBUNIT: DIAGNOSTIC AND TITLE OF INVENTION: THERAPEUTIC METHODS
TITLE OF INVENTION: THERAPEUTIC METHODS
COURSESSANDENCES: 335
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5.8%; Score 26; DB 1; Length 26; 100.0%; Pred. No. 40; cive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTE: AND EMADICAL CELLET, SUIL FLOOR
CONTRY: United States of America
CONTRY: United States of America
ZIP: 94111
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PASTENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/912,951
FILING DATE: 14-AUG-1997
CLASSIFICATION: 435
PRIOR APPLICATION NUMBER: US 08/854,050
FILING DATE: 09-MAY-1997
CLASSIFICATION NUMBER: US 08/851,843
FILING DATE: US 08/851,843
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, 8th Floor
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNDBER: US 08/844,419
FILING DATE: UB-APR-1997
CLASSIFICATION: 435
PRIOR APPLICATION WASE: US 08/724,643
FILING DATE: 01-0CT-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Apple, Randolph T.
REGISTRATION UNDBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002600US
TELEFONE: (415) 576-0200
TELEFONE: (415) 576-0300
TELEFAX: (415) 576-0300
TELEFAX: (415) 576-0300
TELEFAX: (415) 576-0300
TELERATICON FOR SEQ ID NO: 311:
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
TYPE: MUCLEIC acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TCTAACCCTAACTGAGAAGGGCGTAG 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 rcraacccraacrgagagggggrag 26
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Best Local Similarity 100.0
Matches 26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            , MOLECULE TYPE: DNA US-08-912-951-311
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 63
US-08-912-951-312/c
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                                                                                                                                                                                                                                                                                                                    Sequence 4, Application US/09633573

Sequence 4, Application US/09633573

Patent No. 6607898

GENERAL INFORMATION:
APPLICANT: Kopreski, Michael S.
APPLICANT: Gocke, Christopher D.
TITLE OF INVENTION: Telomerase-associated RNA in Plasma or Serum
FILE REFERENCE: 00-1328

CURRENT APPLICATION NUMBER: US/09/653,573

CURRENT FILING DATE: 2000-08-31

PRIOR FILING DATE: 2000-08-31

PRIOR FILING DATE: 2000-08-31

NUMBER OF SEQ ID NOS: 7

SEQ ID NO 4

LENGTH: 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Kopreski, Michael S.
APPLICANT: Kopreski, Michael S.
APPLICANT: Gocke, Christopher D.
TITLE OF INVENTION: Method for Detection of hTR and hTERT
TITLE OF INVENTION: Telomerase-associated RNA in Plasma or Serum
FILE REFERENCE: 00-1328
CURRENT APPLICATION NUMBER: US/09/653,573
CURRENT PILING DATE: 2000-08-31
PRIOR FILING DATE: 2000-08-31
NUMBER OF SEQ ID NOS: 7
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 5
LENGTH: 26
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                                                                                  DB 1; Length 26; 40;
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                                                                                                                               0; Indele
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                  5.8%; Score 26; DB 100.0%; Pred. No. 40; Live 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 5.8%; Score 26; DB Best Local Similarity 100.0%; Pred. No. 40; Matches 26; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5.8%; Score 26; DB
100.0%; Pred. No. 40;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                           45 TCTAACCCTAACTGAGAAGGGCGTAG 70
                                                                                                                                                                                                  1 TCTAACCCTAACTGAGAAGGCGTAG 26
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; Sequence 5. Application US/09653573
; Patent No. 6607998
; GENERAL INFORMATION:
                                                                                  Query Match
Best Local Similarity 100.0
Matches 26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 100.0
Matches 26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          , ORGANISM: Homo sapiens
US-09-653-573-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-653-573-5
linear
E: DNA
TOPOLOGY: 11
MOLECULE TYPE:
                                           US-09-057-351-30
                                                                                                                                                                                                                                                                                  RESULT 66
US-09-653-573-4
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ZIP: 94111-3314

ZIP: 94111-3315

ZIP: 94111-3324

COMPUTER RADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351

FLING DATE: 08-APR-1994
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/272,102

FILING DATE: 07-UL-1994
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-JUN-1995
ATPORNEY/AGENT INFORMATION:
APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-JUN-1995
ATPORNEY/AGENT INFORMATION:
ARBERENCE/DOCKET NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 3576-0300
INFORMATION FOR SEQ ID NO: 30:
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 30:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 30, Application US/09057351
; Sequence 30, Application US/09057351
; Patent No. 6548298
; GENERAL INFORMATION:
; APPLICANT: Villeponteau, Bryant
; APPLICANT: Funk, Walter
; APPLICANT: Andrews, William H.
; TITLE OF INVENTION: Mammalian Telomerase
; CORRESPONDENCES: 42
; CORRESPONDENCES: 42
; CORRESPONDENCES: Townseend and Townseend and Crew Lide
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
COUNTRY: USA
                                                             015389-000821US
                                                                                                                                                                                                                                                                                                                                                    5.8%; Score 26; DB 100.0%; Pred. No. 40; cive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                        145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26 CTTCCACCGTTCATTCTAGAGCAAAC 1
         ATTOKNET AGENT INFORMATION

NAME: Storella, John R.

REGISTRATION NUMBER: 32,944

REFERENCE/OCKET NUMBER: 01536

TELECOMMUNICATION INFORMATION:

TELEPAK: (415) 576-0200

TELEPAK: (415) 576-0300

INFORMATION FOR SEQ ID NO: 23:

SEQUENCE CHARACTERISTICS:

LENGTH: 26 base pairs

TYPE: nucleic acid

STRANDENNESS: single
                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 100.
Matches 26; Conservative
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                                                                                                                                                                                                                                                           ) TOPOLOGY: linear
) MOLECULE TYPE: DNA
US-09-057-351-23
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Sequence 597, Application US/09402181B
Fatent No. 6610839
GENERAL INFORMATION:
Lingner, Joachim
Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin B.
Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 633
CORRESPONDENCE ADRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 9411.3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: 015389-002620US
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SEQUENCE DESCRIPTION: SEQ ID NO: 597:
US-09-402-181B-597
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 14-AUG-1997
APPLICATION WUNBER: US 08/912,951
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-OCT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/724,643

FILING DATE: 01-07-1996

APPLICATION NUMBER: US 08/844,419

FILING DATE: 18-APR-1997

APPLICATION NUMBER: US 08/846,017

FILING DATE: 25-APR-1997

APPLICATION NUMBER: US 08/851,843

FILING DATE: 06-MAY-1997

APPLICATION NUMBER: US 08/851,343

FILING DATE: 08-MAY-1997

APPLICATION NUMBER: US 08/854,050

FILING DATE: 09-MAY-1997

APPLICATION NUMBER: US 08/854,050
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/402,181B
FILING DATE: 29-Sep-1997
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Ausenhus, Scott L.
REGISTRATION NUMBER: 42,271
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPHAY: (415) 576-0300
INFORMATION FOR SEQ ID NO: 597:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY:
RESULT 68
US-09-402-181B-597
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5.8%; Score 26; DB 1; Length 26;

Query Match

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Gaps
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0
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TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 633
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                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: 
CLASSIFICATION: 
CLASSIFICATION: 
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-0CT-1996
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/81,843
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/81,843
FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/911,312
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 01-0CT-1997
TUDORURY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Ausenhus, Scott L.
REGISTRATION NUMBER: 42,271
REFERENCE POCKET NUMBER: 015389-002620US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                  ·
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    Pred. No. 40; ; Mismatches
                                                                                                                                     70
                                                                                                                                                                                                                        1 rcraacccraacrgagagggggrag 26
                                                                                                                                 45 TCTAACCCTAACTGAGAAGGGCGTAG
                                                                                                                                                                                                                                                                                                                                                                                              US-09-402-181B-598/C; Sequence 598, Application US/09402181B
: Patent No. 6610839; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Cech, Thomas R.
Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin B.
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Best Local Similarity 100.0\$_i p_1
Matches 26_i Conservative 0_i
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: 598:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (415)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY:
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                                              Matches
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MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Ploppy disk
COMPUTER: LIBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/09/721,456
FILING DATE: 22-No. 6617110-2000
CIGASIFICATION: «Unknown»
PRIOR APPLICATION NUMBER: US/08/974,549A
FILING DATE: 19-NOV-1997
APPLICATION NUMBER: US 08/724,643
FILING DATE: 10-OCT-1996
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 2-APR-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 2-APR-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 2-APR-1997
APPLICATION NUMBER: US 08/851,843
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Morin, Gregg B.
Harley, Calvin B.
Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 727
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                 Length 26;
                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Townsend and Crew LLP
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                                                                                                                                                                                                                                              NAME/KEY: -
LOCATION: 1..26
OTHER INFORMATION: /note= "F3b primer"
SEQUENCE DESCRIPTION: SEQ ID NO: 597:
                                                                                                                                                                                                                                                                                                                                                                                 DB 1;
40;
                                                                                                                                                                                                                                                                                                                                                           5.8%; Scor.
100.0%; Pred. No. ...
0; Mismatches
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APPLICATION NUMBER: US 08/911,312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US 08/854,050
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 rcraacccraacrgagaaggggrag 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-721-456-598/c
; Sequence 598, Application US/09721456
; Patent No. 6617110
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 06-MAY-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cech, Thomas R.
Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Townsend and
                                                                                                LENGTH: 26 base pairs
                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
                                            INFORMATION FOR SEQ ID NO: 597
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: California
                                                                                                                                                                     TOPOLOGY: linear
MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Cech,
                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 26; Conserv
                                                                                                                                                                                                                                       NAME/KEY:
                      TELEFAX:
                                                                                                                                                                                                                                                                                                                                     US-09-721-456-597
                                                                                                                                                                                                                   FEATURE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 71
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ZIALE: 19411-3834
COMPUTER READABLE FORM:
MEDIUW TYPE: Ploppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATIBLE
COMPUTER: DATA: PC-DOS/MS-DOS
SOFTWARE: PATENTIN PC-DOS/MS-DOS
CURRENT APPLICATION DATA:
APPLICATION DATA: 109/721,456
FILING DATE: 22-No. 6617110-2000
CLASSIFICATION: «UNknown»
PRIOR APPLICATION DATA: APPLICATION DATA: APPLICATION NUMBER: US 08/74,549A
FILING DATE: 19-NOV-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: 10-OCT-1996
APPLICATION NUMBER: US 08/846,017
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 23-APR-1997
APPLICATION NUMBER: US 08/851,843
APPLICATION NUMBER: US 08/851,843
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chapman, Karen B.
Morin, Gregg B.
Harley, Callvin B.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 727
                                                                                                                                            Length 26;
                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Townsend and Townsend and Crew LL STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REGISTRATION INMBER: 36,429
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/854,050
FILING DATE: 09-MAY-1997
APPLICATION NUMBER: US 08/911,312
FILING DATE: 14-AUG-1997
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
FILING DATE: 01-OCT-1997
APPLICATION NUMBER: WO PCT/US97/17618
FILING DATE: 01-OCT-1997
APPLICATION NUMBER: WO PCT/US97/17885
LOCATION: 1..26

COTHER INFORMATION: /note= "R3c primer"

SEQUENCE DESCRIPTION: SEQ ID NO: 598:
US-09-402-181B-598
                                                                                                                                         Query Match 5.8%; Score 26; DB 1;
Best Local Similarity 100.0%; Pred. No. 40;
Matches 26; Conservative 0; Mismatches
                                                                                                                                                                                                                                       145 CTTCCACCGTTCATTCTAGAGCAAAC 170
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Sequence 597, Application US/09721456

Patent No. 6617110

GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cech, Thomas R.
Lingner, Joachim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nakamura, Toru
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Cech,
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Query Match
Best Local Similarity 72.03
Matches 18; Conservative
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STRANDEDNESS: single
nucleic acid
                                    linear
                                                                            ) DESCRIPTION:
) DESCRIPTION:
) DESCRIPTION:
) DESCRIPTION:
US-08-630-019A-26
TYPE: nucleic
STRANDEDNESS:
TOPOLOGY: lin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
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Sequence 26, Application US/08630019A

Patent No. 6015710

GENERAL INFORMATION:
APPLICANT: Wright, Woodring E.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Orey, David
APPLICANT: No. 6015710ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by
TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/630,019A
FILING DATE: 09-UNN-1996
CLASSIFICATION: 536
                                                       ATTORNEY/AGENT INFORMATION:
NAME: APPIG. RANGOLDh TGG
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002610US
TELECOMMUNICATION INFORMATION:
TELEFONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 598:
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDENESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "R3c primer"
                     FILING DATE: 01-OCT-1997
APPLICATION NUMBER: WO PCT/US97/17885
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REPERENCE/DOCKET NUMBER: 015389-001600US
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0300
INPORMATION POR SEQ ID NO: 26: SEQUENCE CHARACTERISTICS:
  APPLICATION NUMBER: WO PCT/US97/17618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5.8%; Score 26; DB 1;
100.0%; Pred. No. 40;
:ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: -
LOCATION: 1..26
COTHER INFORMATION: /note= "R3c;
SEQUENCE DESCRIPTION: SEQ ID NO: 598:
US-09-721-456-598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTTCCACCGTTCATTCTAGAGCAAAC 1
                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 100.0
Matches 26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25 base pairs
                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       USA
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US-08-630-019A-26
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MOLECULE TYPE: other nucleic acid (PNA),
DESCRIPTION: /desc = "peptide nucleic acid (PNA),
DESCRIPTION: where (deoxy)ribose-phosphate linkages are replaced by
DESCRIPTION: where (deoxy)ribose-phosphate linkages are replaced by
DESCRIPTION: N-(2-aminoethy/1)afycine units linked to nucleotide bases via
DESCRIPTION: glycine amino nitrogen through a methylenecarbonyl linker"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 36, Application US/08630019A
Fatent No. 6015710
GENERAL INFORMATION:
APPLICANT: Shay, Jerry W.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Ocrey, David
APPLICANT: No. 6015710ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by
TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 46
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 25; DB 1; Length 25; Pred. No. 45;
                                                                                                                                                                                    Query Match 5.5%; Score 25; DB 1; Length 25; Best Local Similarity 72.0%; Pred. No. 45; Matches 18; Conservative 7; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: ELB PC compatible

COMPUTER: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION UNMBER: US/08/630,019A

FILING DATE: 09-JUN-1996

CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  015389-001600US
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                                                                                                                                                                                                                                                                                                      41 TTTGTCTAACCCTAACTGAGAAGGG 65
                                                                                                                                                                                                                                                                                                                                     1 UUUGUCUAACCCUAACUGAGAAGG 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION UNDRER: 32,94
REFERENCE/DOCKET UNDRER: 01538
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
INFORMATION FOR SEQ ID NO: 36:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: RNA (genomic)
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Sequence 28, Application US/08482115B
Sequence 28, Application:
APPLICANT: Villeporteau, Bryant
APPLICANT: Punk, Walter
APPLICANT: Andrews, William H.
STITLE OF INVENTION: Telomerase
NUMBER OF SEQUENCES: 40
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
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OUTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/08/482,115B
FILING DATE: 07-JUN-1995
CLASSIFICATION: 435
PRIOR APPLICATION: 135
                                                        ZIP: 9411-3834
COMPUTER REALABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/349,532
                                                                                                                                                                                                                                 APPLICATION NOMBER: US/09/349,532
CLASSIFICATION DATE:
CLASSIFICATION DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELEFENNE: (415) 576-0200
TELEFENNE: (415) 576-0300
TELEFEN: (415) 576-0300
TELEFEX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pred. No. 45; 7; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 5.5%; Score 25; Best Local Similarity 72.0%; Pred. No. 4 Matches 18; Conservative 7; Mismatch
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APPLICATION NUMBER: US 08/272,102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: RNA (genomic) US-09-349-532-40
San Francisco
                       California
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                     Sequence 40, Application US/09349532

Batent No. 6294650

GENERAL INFORMATION:

APPLICANT: Shay, Jerry W.

APPLICANT: Wright, Woodring E.

APPLICANT: Corey, David R.

APPLICANT: Corey, David R.

APPLICANT: Corey, David R.

APPLICANT: No. 6294650ton, James C.

TITLE OF INVENTION: Poptide Nucleic Acids

NUMBER OF SEQUENCES: 60

CORRESPONDENCE ADDRESS:

CORRESPONDENCE ADDRESS:

ADDRESSEE: Townsend and Townsend and Crew LLP

STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1; Length 25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      41 TTTGTCTAACCCTAACTGAGAAGGG 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 UUUGUCUAACCCUAACUGAGAAGGG 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: RNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 72.0°
Matches 18; Conservative
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US-09-349-532-40
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Pred. No. 52;
0; Mismatches 0; Indels
                                                                                                      DB 1; Length 28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/057,351
FILNG DATE: 08-APR-1994
CLASSIFICATION NUMBER: US/09/057,351
FILNG DATE: 08-APR-1994
PRIOR APPLICATION NUMBER: US/09/057,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION NUMBER: US/08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US/08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US/08/330,123
FILING DATE: 07-JUN-1995
ATTONNEY/AGENT: INFORMATION:
AND AND APPLICATION NUMBER: US/08/472,802
FILING DATE: 07-JUN-1995
ATTONNEY/AGENT: INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Enbarcadero Center, Eighth Floor CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                        Sequence 29, Application US/09057351
Fatent No. 6548298
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junia
APPLICANT: Funk, Walter
APPLICANT: Funk, Walter
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           015389-000821US
                                                                                                      Query Match 5.5%; Score 25; DB Best Local Similarity 100.0%; Pred. No. 52; Matches 25; Conservative 0; Mismatches
                                                                                                                                                                                                   81 TTTGCTCCCCGCGCGCTGTTTTCT 105
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CONTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
WEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC_DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5.5%; Scc.
100.0%; Pre
0; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFRENCE/DOCKET NUMBER: 01531
TELECOMMUNICATION INFORMATION:
TELEPAX: (415) 576-0200
INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 100.
Matches 25; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 28 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
            TOPOLOGY: linear
                                  MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                          US-09-057-351-29/c
                                              , WS-08-472-802C-29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1; Length 28; 52;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OPERATING SYSTEM: PC-UDS/MS-LDS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/472,802C
FILING DATE: 07-JUN-1995
CLASSIFACATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
FILING DATE: 07-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAMM: SMICH, William M.
REGISFRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 15389-000820
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INCORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Villeponteau, Bryant
APPLICANT: Peng, Junii
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 4
NUMBER OF SEQUENCES: 4
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
FILING DATE: 07-JUL-1994

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
ATTORNEY/AGENT INFORMATION:
NAME: SECOFILA, JOHN R.
REGISTRATION NUMBER: 015389-000830US
TELECOWNUNICATION INFORMATION:
TELEPHONE: (415) 576-030
TELEPAX: (415) 676-030
TELENGHH: 28 base pairs
LENGHH: 28 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5.5%; Score 25; DB 100.0%; Pred. No. 52; ive 0; Mismatches
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Sequence 29, Application US/08472802C
; Patent No. 5958680
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEO ID NO: 29
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 28 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    USA
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US-08-472-802C-29/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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5.3%; Score 24; DB 1; Length 24;
100.0%; Pred. No. 52;
tive 0; Mismatches 0; Indels
                                                                                                                                    MEDIUM ITEE: FIODPY GIBE
COMPUTER: IBM PC compatible
SOCTHARE: Patentin Release #1.0, Version #1.30
SOCTHARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INPORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 015389-001610US
TELECHONGION INFORMATION:
TELECHO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: PATENTIN RALease #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/349,532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     41 TTTGTCTAACCCTAACTGAGAAGG 64
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           other nucleic acid
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                            ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 100. Matches 24; Conservative
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
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MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DESCRIPTION
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US-09-349-532-25
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COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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Patent No. 6046307

GENERAL INFORMATION:
APPLICANT: Wright, Woodring E.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Corey, David R.
APPLICANT: No. 6046307, Con. James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by
TITLE OF INVENTION: Peptide Nucleic Acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER IS Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/770,565
FLING DATE: 20-DEC-1996
CLASSIFICATION: 435
ATTORNEY AGENT INFORMATION:
NAME: Storella, John R.
REFERENCE/DOCKET NUMBER: 015389-002300US
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSE: Townsend and Townsend and Crew LLP
STEET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5.4%; Score 24.4; D
96.2%; Pred. No. 62;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           355 CCTTTCAGGCCGCAGGAAGAGGAACG 380
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Best Local Similarity 96.2
Matches 25; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             California
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US-08-770-565-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
                                                                      US-08-770-565-2/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ò
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/desc = "peptide nucleic acid (PNA), where (deoxy(ribose-phosphate linkages are replaced by N-(2-aminoethyl)glycine units linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker" ö Gaps ö APPLICANT: Wight, Woodring E. APPLICANT: Wight, Woodring E. APPLICANT: Wight, Woodring E. APPLICANT: Wistyszek, Mieczyslaw A. APPLICANT: Corey, David R. APPLICANT: Corey, David R. Corestylon: Modulation of Mammalian Telomerage by TITLE OF INVENTION: Modulation of Mammalian Telomerage by TITLE OF INVENTION: Peptide Nucleic Acids NUMBER OF SEQUENCES: 60 CORRESPONDENCE ADDRESS: ADDRESSER: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California

```
APPLICANT: Silverman, Robert H.
APPLICANT: Silverman, Robert H.
APPLICANT: Kondo, Seiji
APPLICANT: Kondo, Seiji
APPLICANT: Cowell, John K.
APPLICANT: Li, Guiying
APPLICANT: Torrence, Paul F.
TITLE OF INVENTION: RAASE L ACTIVATORS AND ANTISENSE OLIGONUCLEOTIDES
TITLE OF INVENTION: EPECTIVE TO TREAT TELOMERASE-EXPRESSING MALIGNANCIES
FILE REFERENCE: 8656-022
CURRENT FILLING DATE: 1999-02-03
EARLIER APPLICATION NUMBER: 60/044,507
EARLIER PILING DATE: 1997-04-21
NUMBER OF SEQ ID NOS: 9
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO S
LENGTH: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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Sequence 26, Application US/08838545

Sequence 26, Application US/08838545

Patent No. 6046307

GENERAL INFORMATION:

APPLICANT: Wisight, Woodring E.

APPLICANT: Pratyszek, Mieczyslaw A.

APPLICANT: Orey, David R.

APPLICANT: No. 6046307ton, James C.

TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Modulation of Mammalian Telomerase by CORRESPONDENCE ADDRESS:

CORRESPONDENCE ADDRESS:

ADDRESSEE: Two Embarcadero Center, Eighth Floor STATE: California

COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ·.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: Description of Artificial Sequence: primer US-09-018-125-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 5.3%; Score 24; DB 1; Length 24; Best Local Similarity 100.0%; Pred. No. 52; Matches 24; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SIKELL
CITY: San Frantic.
STATE: California
STATE: California

ZIP: 9411.3334

ZIP: 9411.3334

COUNTRY: USA

ZIP: 9411.3334

COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PatentIn Release #1.0, Version #1.30

SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/838,545

FILING DATE: 09-APR-1997

CIASSIFICATION NUMBER: US 08/630,019

FILING DATE: 09-APR-1996

ATTORNEY AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944

REFERRNCE/DOCKET NUMBER: 015389-001610US

TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200

(415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           423 CGTGCACCCAGGACTCGGCTCACA 446
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ð
               ATTORNES' 09-APR-1997

APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996

ATTORNES' AGENT INFORMATION:
NAME: Storella John R.
FREGISTRATION NUMBER: 12,944

REGISTRATION NUMBER: 12,944

REFERENCE/DOCKET NUMBER: 015389-001610US

TELEPHONE: (415) 576-0200

TELEPHONE: (415) 576-0200

TELEPAX: (415) 576-0300

INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
LENGTH: 24 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
SECRIPTION: (desc. "peptide nucleic acid (FNA), DESCRIPTION: where (decxy(ribose-phosphace linkages are replaced by DESCRIPTION: Qestallary) glycine units linked to nucleotide bases via
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Silverman, Robert H.
APPLICANT: Kondo, Seiji
APPLICANT: Kondo, Seiji
APPLICANT: Modo, Seiji
APPLICANT: Modo, Seiji
APPLICANT: O'Gulying
APPLICANT: Li, Gulying
APPLICANT: Torrence, Paul F.
TITLE OF INVENTION: EFFECTIVE TO TREAT TELOMERASE-EXPRESSING MALIGNANCIES
TITLE OF INVENTION: EFFECTIVE TO TREAT TELOMERASE-EXPRESSING MALIGNANCIES
TITLE OF INVENTION: EFFECTIVE TO TREAT TELOMERASE-EXPRESSING MALIGNANCIES
TITLE OF INVENTION: 1999-02-03
CURRENT APPLICATION NUMBER: 09902-03
CURRENT APPLICATION NUMBER: 60/044,507
EARLIER PILING DATE: 1997-04-21
MUMBER OF SEQ ID NOS: 9
SOFTWARE: PATENTIN VOIL 2.0
SEQ ID NO 4
ENGINE PATENTIAL VOIL 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 5.3%; Score 24; DB 1; Length 24; Best Local Similarity 100.0%; Pred. No. 52; Matches 24; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5.3%; Score 24; DB 1; Length 24;
100.0%; Pred. No. 52;
Live 0; Mismatches 0; Indels
   US 08/838,545
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 TTGTCTAACCCTAACTGAGAAGG 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Sequence 4, Application US/09018125A; Patent No. 6468983
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 83
US-09-018-125-5/c
Squence 5, Application US/09018125A
Patent No. 6466983
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 100.0
Matches 24; Conservative
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-349-532-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-018-125-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   JS-09-018-125-4
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US-08-485-778-20
                                                                                                                                                                                                                        RESULT 86
US-08-485-778-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                  요
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/desc = "peptide nucleic acid (PNA),
where (descytibose-phosphate linkages are replaced by
N.Cz-aminoethyl jlycine units linked to nucleotide bases via
glycine amino N through a methylenecarbonyl linker
                                                                                                                                                       bases via
                                                                                                                                                                                                                                                                                         ö
                                                                                   other nucleic acid
/desc = "peptide nucleic acid (PNA),
where (deoxy(ribose-phosphate linkages are replaced by
N-(2-aminoethyl)glycine units linked to nucleotide base
glycine amino N through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Shay, Jerry W.
APPLICANT: Wright, Woodring E.
APPLICANT: Piatyasek, Mieczyslaw A.
APPLICANT: Corey, David R.
APPLICANT: No. 6294650ton, James C.
APPLICANT: No. 6294650ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                        5.1%; Score 23; DB 1; Length 23; 100.0%; Pred. No. 58; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 94111-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/349,532
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         015389-001610US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION:
PRIOR APPLICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-0016:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFAX: (415) 576-0200
INFORMATION FOR SEQ ID NO: 26:
SEGUIENCE CHARACTERISTICS:
LENGTH: 23 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                               1 CCATTTTTGTCTAACCCTAACT 23
                                                                                                                                                                                                                                                                                                                                     35 CCATTTTTGTCTAACCCTAACT 57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 26, Application US/09349532
Patent No. 6294650
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                         23; Conservative
23 base pairs
                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          single
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Best Local Similarity
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                                                                 linear
                                                            TOPOLOGY: line
MOLECULE TYPE: C
DESCRIPTION: /
DESCRIPTION: W
DESCRIPTION: W
DESCRIPTION: W
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DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
US-09-349-532-26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-349-532-26
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  DB 1; Length 23;
58;
                                                Indels
                                                                                                                                                                                                                                                                                                                    APPLICANT: Avilion, Ariel Athena
APPLICANT: Avilion, Ariel Athena
APPLICANT: Feng, Junia
APPLICANT: Feng, Junia
APPLICANT: Funk, Walter
APPLICANT: Greider, Carol
APPLICANT: Marhuenda, Maria Antonia Blasco
APPLICANT: Willeponteau, Bryant
TITLE OF INVENTION: NA COMPONENT OF TELOWERASE
NUMBER OF SEQUENCES: 45
CORRESPEDINGE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: Two Militia Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5.1%; Scor.
100.0%; Pred. No. ...
Query Match 5.1%; Score 23; DB Best Local Similarity 100.0%; Pred. No. 58; Matches 23; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRENT A PERILCATION DATA:
CLASSIPECATION TOTAS:
FILING DATE: 07-JE-1995
CLASSIPECATION 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/387,524
FILING DATE: 13-FEB-1995
FILING DATE: 13-FEB-1995
FILING DATE: 27-OCT-1994
FILING DATE: 07-JUL-1994
ATORNEY/AGENT INFORMATION:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
ATORNEY/AGENT INFORMATION:
ARGISTRATION NUMBER: 32,227
REFERENCE/DOCKET NUMBER: 32,227
FELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                            35 CCATTTTTGTCTAACCCTAACT 57
                                                                                                                              1 CCATTTTTTGTCTAACCCTAACT 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   44
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                                                                                                                                                                                                                                                       ; Sequence 20, Application US/08485778; Patent No. 5876979
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 02173
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
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Best Local Similarity
Matches 23; Conservat
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4.9%; Score 22; DB 1; Length 22; 100.0%; Pred. No. 66; 0; Indels tive 0; Mismatches 0; Indels
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APPLICANT: FENG, Juni,
APPLICANT: FENG, Juni,
APPLICANT: FENG, Juni,
APPLICANT: FUNK, Walter
APPLICANT: ADDRESS:
CORRESPONDENCE: 25
CORRESPONDENCE ADDRESS:
ADDRESSE: Townsend and Townsend Khourie and Crew
STREET: 379 Lytton Avenue
CITY: Palo Alto
                                                                                                                                                                               COUNTER: US
ZIN: 94301

COMPUTER READABLE FORM:
MEDIUM TYDE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 27-0CT-1994
CLASSIFICATION NUMBER: US 08/272,102
FILING DATE: 37-0CT-1994
ATTORNEY/AGENT INPORMATION:
PROTESTRAINON NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 15389-000810
TELEPONE: (415) 326-2420
TELEPONE: (415) 326-2420
TELEPONE: (415) 326-2422
SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
TYPE: nucleic acid
TOPOLOGY: Innear
NOTHER TYPE: RNA
                        CORRESPONDENCE ADDRESS:
ADDRESSE: Townsend and Townsend Khourie and Crew STREET: 379 Lytton Avenue
CITY: Palo Alto
STATE: California
COUNTRY: US
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/NS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
FILING DATE: 27-OCT-1994
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  46 CTAACCCTAACTGAGAAGGGCG 67
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; Sequence 6, Application US/08330123A
; Patent No. 5583016
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity 100.
Matches 22; Conservative
NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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                                                                                                        APPLICANT: Andrews, William H.
APPLICANT: Andrews, William H.
APPLICANT: Andrews, William H.
APPLICANT: Peng, Junil
APPLICANT: Marthenda, Maria A. B.
APPLICANT: Marthenda, Maria A. B.
APPLICANT: Marthenda, Bryant
ITILE OF INVENTION: RNA Component of Telomerase
NUMBER OF SEQUENCES: 47
CORRESPONDENCE ADDRESS:
ADDRESSE: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: Two Militia Drive
CITY: Lexington
STREET: Fro Militia Drive
CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: WA
COUNTRY: US
CONTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5.1%; Score 23; DB 1; Length 27; ilarity 100.0%; Pred. No. 70; Conservative 0; Mismatches 0; Indels
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Sequence 5, Application US/08330123A

Patent No. 5583016

GENERAL INFORMATION:
APPLICANT: VILLEPONTEAU, Bryant
APPLICANT: FENG, Junli
APPLICANT: FUNK, Walter;
APPLICANT: FUNK, Walter;
TITLE OF INVENTION: HUMAN TELOMERASE
                                        Sequence 20, Application US/08520550A Patent No. 6013468 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22 GGAGGGGTGGTGGCCATTTTTG 44
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
nes 23; Conserva
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               US-08-520-550A-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; TOPOLOGY:
US-08-520-550A-20
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Query Match
Best Local Similarity 100.0
Matches 22; Conservative
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Best Local Similarity 100.0
Matches 22; Conservative
                                                                                                                                                                                                                                                        183 CTGCTGGCCCGTTCG
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STRANDEDNESS: single
TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-660-678A-27/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 92
US-08-660-678A-28/c
                                                              US-08-482-115B-32
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Sequence 32, Application US/08482115B
Patent No. 576679
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Juni
TAPLICANT: Feng, Juni
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Assays for the RNA Component of Human
TITLE OF INVENTION: Telomerase
NUMBER OF SEQUENCES: 40
CORRESPONDENCE ADDRESS: 40
CORRESPONDENCE ADDRESS: 40
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Townsend and Townsend and Crew LLP
STREET: San Francisco
STATE: California
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Pred. No. 66;
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MEDIUM TYPE: Floppy disk

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC.DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURSTITION DATA:

APPLICATION NUMBER: US/08/482,115B

FILING DATE: 07-JUN-1995

CLASSIFICATION: 435

RIGH APPLICATION DATA:

APPLICATION NUMBER: US 08/272,102

FILING DATE: 07-JUL-1994

PRIOR APPLICATION NUMBER: US 08/330,123

FILING DATE: 27-OCT-1994

PRIOR APPLICATION NUMBER: 32,944

REFERENCE/DOCKET NUMBER: 32,9
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100.0%; Pred. No. v..
0; Mismatches
                       FILING DATE: 07-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: SMITCH, William W
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 15389-000810
TELEPRACIONOMICATION INFORMATION:
TELEPRACIONE: (415) 326-2400
TELEPRACIONE: (415) 326-2420
INFORMATION FOR SEQ ID NO: 6: SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: RNA
   US 08/272,102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            54 AACTGAGAAGGGCGTAGGCGCC 75
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Best Local Similarity 100.
Matches 22; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bingle
   APPLICATION NUMBER:
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STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     USA
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DB 1; Length 22;
66;
                                                  0; Indels
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ZIP: 94111-3834

COUNTRY: USA

ZIP: 94111-3834

COMPUTER READABLE FORM:

MEDIUM TYEE: FORM:

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/660,678A

FILING DATE: 05-JUN-1996

CLASSIFICATION DATA:

APPLICATION NUMBER: US 08/330,123

FILING DATE: 27-OCT-1994

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/22,102

FILING DATE: 07-JUL-1994

ATTORNEY/AGENT INFORMATION:

NAME: SLOCAFILA, DATA:

REGISTRATION NUMBER: 015389-000811US

TELECOMMUNICATION INFORMATION:

NAME: SLOCAFILA, STOCAFORDER: O15389-000811US

TELECOMMUNICATION INFORMATION:

MET DEVICE FORCEST NUMBER: 015389-000811US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Townsend and Townsend and Crew LDI STREET: Two Embarcadero Center, Eighth Floor CITY: San Prancisco STATE: California
  4.9%; Score 22; DB 100.0%; Pred. No. 66; ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                          APPLICANT: French, Junii
APPLICANT: French, Walter
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Cr
                                                                                                  CCTCC 204
                                                                                                                                                                                                                                                               Sequence 27, Application US/08660678A; Patent No. 5837857; GENERAL INFORMATION: APPLICANT: Villeponteau, Bryant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             46 CTAACCCTAACTGAGAAGGGCG 67
                                                                                                                                22 CTGCTGGCCCGTTCGCCCCTCC 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (415) 576-0200
TELEPAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 27:
SEQUENCE CHARACTERISTICS:
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Gaps
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                                                                                                                              COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/485,778
FILING DATE: 07-JE-1995
CLLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/387,524
FILING DATE: 13-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
ATTONNEY/AGENT INFORMATION:
ANAME: O7-JUL-1994
ATTONNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C. STREET: Two Militia Drive CITY: Lexington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Andrews, William H.
APPLICANT: Avilion, Ariel Athena
APPLICANT: Feng, Jun1i
APPLICANT: Feng, Jun1i
APPLICANT: Feng, Jun1i
APPLICANT: Greider, Carol
APPLICANT: Greider, Carol
APPLICANT: Marhuenda, Marria Antonia Blasco
APPLICANT: Willeponteau, Bryant
TITLE OF INVENTION: RNA COMPONENT OF TELOMERASE
NUMBER OF SEQUENCES: 45
CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.(
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PatentIn Release #1.0, Version #1.30
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      46 CTAACCCTAACTGAGAAGGGCG 67
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Patent No. 5876979
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Granahan, Patricia
REGIGTRATION NUMBER: 32,227
REFERENCE/DOCKET NUMBER: CSH
TELECOMMUNICATION INFORMATION:
TELERBHONE: 617-861-6240
TELEFAX: 617-861-9540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                          COMPUTER READABLE FORM:
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US-08-485-778-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 22; Conserv
                                                                                                                     MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: MA
COUNTRY: US
ZIP: 02173
                                                                    02173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-485-778-8/c
                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH:
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Pred. No. 66;
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CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/660,678A
FILING DATE: 05-JUN-1996
CLASSITEATION: 435
FILING DATE: 05-JUN-1996
CLASSITCATION DATA:

APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
FILING DATE: 27-OCT-1994
PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
ATTORNEY/AGENT INFORMATION:

NAME: Storella, John R.

REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000811US
TELECHMAUNICATION INFORMATION:

TELECHMACE ACCOUNTION OF STG-0300
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3: Hamilton, Brook, Smith & Reynolds, P.C.
Two Militia Drive
       Patent No. 597857;
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Julia
APPLICANT: Feng, Julia
APPLICANT: Funk, Walter
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
COUNTRY: USA
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APPLICANT: Funk, Walter
APPLICANT: Greider, Carol
APPLICANT: Greider, Carol
APPLICANT: Villeponteau, Bryant
TITLE OF INVENTION: RNA COMPONENT OF TELOWERASE
CORRESSPONDENCES: 45
CORRESSPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Andrews, William H.
Avilion, Ariel Athena
Feng, Junli
Funk, Walter
Greider, Carol
Marhuenda, Maria Antonia Blasco
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                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      54 AACTGAGAAGGGCGTAGGCGCC 75
Sequence 28, Application US/086606781
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 100.u
....hes 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear
MOLECULE TYPE: DNA
US-08-660-678A-28
                                                                                                                                                                                                                                                                                                                                                   94111-3834
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US-08-485-778-7/c
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET:
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100.0%; Pred. No. 66;
cive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                             DB 1; Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
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SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/472,802C
FILING DATE: 07-JUN-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 07-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAMME: SMith, William M.
REGISTRATION NUMBER: 15389-000820
TELEPOMOKET NUMBER: 15389-000820
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 42:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 42, Application US/08472802C
Sequence 42, Application US/08472802C
Patent No. 5958680
GENERAL INFORMATION:
APPLICANT: Feng, Juni;
APPLICANT: Feng, Juni;
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSER: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                      4.9%; Score 22; DB 100.0%; Pred. No. 66; tive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             46 CTAACCCTAACTGAGAAGGGCG 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CTGCTGGCCCGTTCGCCCCTCC 1
                            TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 37
SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 100.0
Matches 22, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 100.0
Matches 22; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                          linear
                                                                                                                                                                                                                                      ; TOPOLOGY: lii
; MOLECULE TYPE:
US-08-472-802C-37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: 11 MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-472-802C-42/c
   TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4.9%; Score 22; DB 1; Length 22;
100.0%; Pred. No. 66;
iive 0; Mismatches 0; Indels
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MEDIUM TYPE: Floppy disk
COMPUTER: ISPA
COMPUTER: ENDORY disk
COMPUTER: ENDORY disk
COMPUTER: ISPA
COMPOTER: OF COMPATIBLE
COMPOTER: OF COMPATIBLE
COMPOTER: OF COMPATIBLE
SUSTINGENT APPLICATION DATA:
APPLICATION NUMBER: US/08/472,802C
FILING DATE: O7 -JUN-1995
CLLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-CT-1994
ATORING DATE: 27-CT-1994
ATORING DATE: 37-CT-1994
ATORING WARE: MILITIAM M.
REGISTRATION NUMBER: 30,223
REGISTRATION NUMBER: 30,223
REGISTRATION NUMBER: 30,223
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| Patent No. 5958680
| GENERAL INFORMATION:
| APPLICANT: Villeponteau, Bryant
| APPLICANT: Feng, Juni |
| APPLICANT: Andrews, William H. |
| TITLE OF INVENTION: Mammalian Telomerase |
| TITLE OF INVENTION: 44 |
| CORRESPONDENCE ADDRESS: 44 |
| ADDRESSEE: Townsend and Townsend and Crew LLP |
| STREET: Two Embarcadero Center, Bighth Floor |
| CITY: San Francisco |
| CITY: San Francisco |
| CONTRESS |
| CONTRESSEE: Contact |
| CONTRESSEE: Townsend and Townsend and Crew LLP |
| CONTRESSEE: Townsend and Townsend and Crew LLP |
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| CONTRESSEE: Townsend and Townsend and Crew LLP |
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FELECOMMUNICATION INFORMATION:
                            FILING DATE: 07-JE-1995
CLASSIPTICATION: 435
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
FILING DATE: 13-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/22,102
FILING DATE: 07-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Granahan, Patricia
RECISTRATION NUMBER: 32,227
RECCOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CSHL94-05A4
UMBER: US/08/485,778
07-JE-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 8: SEQUENCE CHARACTERISTICS: LENGTH: 22 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 100.
Matches 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: single
   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-08-472-802C-37/c
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STATE: Ca
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4.9%; Score 22; DB 1; Length 22; 100.0%; Pred. No. 66;
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MEDLIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURENT APPLICATION DATA:
APPLICATION NUMBER: US/08/520,550A
FILING DATE: 29-AUG-1995
CLASSIFICATION NUMBER: US 08/387,524
FILING DATE: 13-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: TO-TUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: O7-UUL-1994
PRIOR APPLICATION NUMBER: US 08/272,102
FILING DATE: O7-UUL-1994
ATTORNEY/AGENT INFORMATION:
AAPLICATION NUMBER: 32,227
REGISTRATION NUMBER: 32,227
RECOMMUNICATION NUMBER: CSHL94-05A3B
TELEPHONE: 617-861-6240
INFORMATION FOR SEQ ID NO: 7: SEQUENCE CHARACTERISTICS:
LEMOTH: 22 base pairs
LEMOTH: 22 base pairs
TELEPHONES: SINGle
TYPE: UNCLEIC acide
STRANDEDNESS: SINGle
TITLE OF INVENTION: RNA Component of Telomerase NUMBER OF SEQUENCES: 47
CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C. STREET: Two Militia Drive
CITY: Lexington
STATE: MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C. STREET: Two Militia Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Andrews, William H.
APPLICANT: Andrews, William H.
APPLICANT: Feng, Junli
APPLICANT: Fronk, Walter
APPLICANT: Fronk, Walter
APPLICANT: Fronk, Walter
APPLICANT: Marhuenda, Maria A. B.
APPLICANT: Willeponceau, Bryant
TITLE OF INVENTON: RNA COMPONENT OF SEQUENCES: 47
CORRESPONDENCES ADRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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; Sequence 8, Application US/08520550A
; Patent No. 6013468
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100.0%; Pit
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nes 22; Conservative
                                                                                                                                                           COUNTRY: US
ZIP: 02173
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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Matches
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                                                                                                                                                                                                                                                                                                                                                                          CITY: CITY: Califor...

STATE: Califor...

COMPUTER: BADABLE FORM:

MBDIUM TYPE: Flopy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PAFCHIIN BATA:

APPLICATION DATA:

APPLICATION NUMBER: US/08/472,802C

FILING DATE: 07-JUN-1995

CLASSIFICATION DATA:

APPLICATION NUMBER: US 08/272,102

FILING DATE: 07-JUN-1994

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/330,123

FILING DATE: 27-OCT-1994

ATTORNEY/AGENT INPORMATION:

NAME: Smith, William M.

REGISTRATION NUMBER: 30,223

REGISTRATION NUMBER: 35,70-020

TELECOMMUNICATION INFORMATION:

TELEPHONE: (415) 576-0300

TELEPHONE: (415) 576-0300

TELEPHONE: (415) 576-0300

INFORMATION FOR SEC ID NO: 43:

SEQUENCE CHARACTERISTICS:

'ENGTH: 22 base pairs

'ENGTH: 22 base pairs

TOTOME ACTION INCLES:

'ENGTH: 22 base pairs
                                                                                                      US-08-472-802C-43/c

| Sequence 43, Application US/08472802C
| Patent No. 5958880
| Patent No. 1958880
| Patent No. 1958880|
| APPLICANT: Villeponteau, Bryant
| APPLICANT: Peng, Junii
| APPLICANT: Andrews, William H.
| TITLE OF INVENTION: Mammalian Telomerase
| NUMBER OF SEUENCES: 44
| CORRESPONDENCE ADDRESS: 44
| CORRESPONDENCE ADDRESS: 57RET: Two Embarcadero Center, Eighth Floor
| CITY: San Francisco
| STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 7, Application US/08520550A
Fatent No. 6013468
Fatent No. 6013468
FAPLICANT: Andrews, William H.
APPLICANT: Avilion, Ariel A.
APPLICANT: Peng, Junii
APPLICANT: Peng, Junii
APPLICANT: Peng, Amiter
APPLICANT: Marhuenda, Maria A. B.
APPLICANT: Marhuenda, Maria A. B.
APPLICANT: Willeponteau, Bryant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Andrews, William H.
Avillon, Axiel A.
Feng, Junii
Funk, Walter
Greider, Carol
Marhuenda, Maria A. B.
Villeponteau, Bryant
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                22 craacccraacrgagagggg 1
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Best Local Similarity
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US-08-520-550A-7/c
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patemetin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/998,443
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNET AND ATTORNET AND ATTORNET AND AND R.
NAME: Stochalla, John R.
REFERENCE/DOCKET NUMBER: 015389-000811US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
       PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-011-1994
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 015389-000811US
TELECOMMUNICATION INPORMATION:
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 27:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-1998-443-28/C
; Sequence 28, Application US/08998443
; Patent No. 6054575
; GENERAL INFORMATION:
; APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junii
; APPLICANT: Funk, Walter
; APPLICANT: Andrews, William H.
; TITLE OF INVENTION: Mammalian Telomerase
; NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRICR APPLICATION DATA:
APPLICATION NUMBER: US/08/660,678
FILING DATE: 05-UN-1996
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRICR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-ULL-1994
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 100.0
Matches 22, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   single
                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
CLASSIFICATION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/520,550A
FILING DATE: 29-ANG-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/387,524
FILING DATE: 11-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-AUL-1994
ATYORNEY/AGENT INFORMATION:
NAME: GRANABATION:
NAME: GRANABATION
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 27, Application US/08998443
; Sequence 27, Application US/08998443
; Patent No. 6054575
; GENERAL INFORMATION:
    APPLICANT: Feng, Junii
    APPLICANT: Feng, Junii
    APPLICANT: Feng, Junii
    APPLICANT: Punk, Walter
    APPLICANT: Nonerage and Townsend and Crew LLP
    STREET: Townsend and Townsend and Townsend
    STREET: Townsend and Townsend

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OPERATING SYSTEM: PC-DOS/MS-DOS
SOTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/998,443
FILING DATE:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/660,678
FILING DATE: 05-JUN-1996
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 617-861-9540
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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Best Local Similarity 100.C
Matches 22; Conservative
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US-08-520-550A-8
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Sequence 6, Application US/09580517
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MEDIUM TYPER: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/580.517

FILING DATE: 25-May-2000

CLASSIFICATION: -UNKNOWN->

PRIOR APPLICATION: -UNKNOWN->

PRIOR APPLICATION NUMBER: 08/330,123

APPLICATION NUMBER: 08/330,123

ATTORNEY/AGENT INFORMATION:

NAME: Smith, William M

REGISTRATION NUMBER: 30,223

REFERENCE/DOCKET NUMBER: 15389-000810

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ANDREWS, William H.
TITLE OF INVENTION: HUMAN TELOMERASE
NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
ADDRESSE: Townsend and Townsend Khourie and Crew
                                                                                                                                        4.9%; Score 22; DB 1; Length 22; 100.0%; Pred. No. 66; 0; Indels tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4.9%; Score 22; DB 1; Length 22;
100.0%; Pred. No. 66;
cive 0; Mismatches 0; Indels
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MOLECULE TYPE: RNA
SEQUENCE DESCRIPTION: SEQ ID NO: 5:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: VILLEPONTEAU, Bryant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 379 Lytton Avenue
CITY: Palo Alto
STATE: California
                                                                                                                                                                                                                                          54 AACTGAGAAGGGCGTAGGCGCC 75
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US-09-580-517-5/c
; Sequence 5, Application US/09580517
; Patent No. 6320039
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (415) 326-2422 INFORMATION FOR SEQ ID NO: 5:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FENG, Junli
FUNK, Walter
                                                                                                                                        Query Match
Best Local Similarity 100.(
Matches 22; Conservative
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Matches 22; Conservative
nucleic acid
                 STRANDEDNESS: sir
TOPOLOGY: linear
MOLECULE TYPE: DNA
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US-08-998-443-28
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RESULT 103 US-09-580-517-6/c

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COUNTRY:
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                                                                                                                                                                                NAME/KEY: misc feature

LOCATION: (1) —

DOTHER INFORMATION: Biotin S'-terminal

OTHER INFORMATION: Description of Artificial Sequence: Affinity Agent

US-09-717-828B-2
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OTHER INFORMATION: Description of Artificial Sequence: Affinity Agent
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4.9%; Score 22; DB 1; Length 22;
Best Local Similarity 100.0%; Pred. No. 66;
Matches 22; Conservative 0; Mismatches 0; Indels
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100.0%; Pred. No. 66;
tive 0; Mismatches 0; Indels
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GRENEAL INFORMATION:

APPLICANT: Atkinson III, Edward M
APPLICANT: Lichtsteiner, Serge P
APPLICANT: Lichtsteiner, Serge P
APPLICANT: Pusarn, Romald A
APPLICANT: Pruzan, Romald A
TITLE OF INVENTION: A Method for Purifying Telomerase
CURRENT APPLICATION NUMBER: US/09/717,829A
CURRENT APPLICATION NUMBER: 09/420,056
PRIOR PELING DATE: 1999-10-18
PRIOR PELING DATE: 1999-10-18
PRIOR PELING DATE: 1995-04-04
PRIOR PILING DATE: 1995-04-04
PRIOR FILING DATE: 1995-08-04
NUMBER OF SEQ ID NOS: 11
SOFTWARE: PATENTIN VUMBER: 06/510,736
PRIOR FILING DATE: 1995-08-04
SEQ ID NO 2
LENGTH: 22
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; Sequence 2, Application US/09717829A
Patent No. 6545133
; GENERAL INFORMATION:
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Patent No. 6548298
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Peng, Junli
APPLICANT: Feng, Junli
                   NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin Ver. 2.1 edited
SEQ ID NO 2
LENGTH: 22
                                                                                                                  TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA ORGANISM: Artificial Sequence
PRIOR FILING DATE: 1995-08-04
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US-09-057-351-41/c
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US-09-717-829A-2
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                                                                                                                                                                  FEATURE:
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Gaps
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66;
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
CLASSIFICATION: 435
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STREET: Two Embarcadero Center, Eighth Floor STATE: California
COUNTRY: USA
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-0TL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
APPLICATION NUMBER: US 08/320,123
FILING DATE: 27-0CT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-0TN-1995
ATTORNEY/AGENT INFORMATION:
NAME: SECREILA, JOHN R.
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
RECECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
INFORMATION FOR SEQ ID NO: 41: SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junii
APPLICANT: Funk, Walter
APPLICANT: Andrews, Walliam H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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; Sequence 42, Application US/09057351
Patent No. 6548298
; GENERAL INFORMATION:
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Best Local Similarity 100.0%; P
Matches 22; Conservative 0;
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: single
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100.0%; Pred. No. 74;
                          4.9%; Score 22; DB 1; Length 22; 100.0%; Pred. No. 66; cive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/330,123A
FLING DATE: 27-CCT-1994
CLASSTETATION NUMBER: US 08/272,102
FILING DATE: O7-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 15389-000810
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION OF 25:
SEQUENCE CHARACTERISTICS:
LUMPOTH: 21 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Townsend and Townsend Khourie and Crew STREET: 379 Lytton Avenue CITY: Palo Alto STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Prec. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INPORMATION:
APPLICANT: VILLEPONTEAU, Bryant
APPLICANT: FENG, Junii
APPLICANT: FUNK, Walter
APPLICANT: ANDREWS, William H.
TITLE OF INVENTION: HUMAN TELOMERASE
NUMBER OF SEGUENCES: 25
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 110
US-08-482-115B-25/C
; Sequence 25, Application US/08482115B
; Patent No. 5776679
; GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junli
; APPLICANT: Funk, Walter
; APPLICANT: Andrews, William H.
                                                                                                                                                                                                                                                                                                                                                                                                  US-08-330-123A-25/c
; Sequence 25, Application US/08330123A
; Patent No. 5583016
                                                                                                                                                                                   46 CTAACCCTAACTGAGAAGGGCG 67
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Best Local Similarity 100.(
Matches 21; Conservative
                                                                                                           22; Conservative
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EDNESS: single
                                  Query Match
Best Local Similarity
Matches 22; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ns
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US-10-330-372-2/C

Patent No. 6787133

GENERAL INFORMATION:

APPLICANT: Weinrich, Scott

APPLICANT: Weinrich, Scott

APPLICANT: Weinrich, Scott

APPLICANT: Weinrich, Scott

APPLICANT: Vasserot, Alain

APPLICANT: Vasserot, Alain

APPLICANT: Vasserot, Alain

TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators and

TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators

TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators

TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators

TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators

TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators

TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators

TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators

FILE REFERENCE: 01/006C

CURRENT FILING DATE: 1995-08-04-04

PRIOR FILING DATE: 1997-04-04

PRIOR FILING DATE: 1999-10-18

PRIOR FILING DATE: 1999-10-18

PRIOR FILING DATE: 1999-10-18

PRIOR FILING DATE: 1999-10-18

PRIOR FILING DATE: 1990-11-20

NUMBER OF SEQ ID NOS: 11

SOUTHARE PATENTION NUMBER: 09/420,056

PRIOR FILING DATE: 1990-11-20

NUMBER OF SEQ ID NOS: 11

SOUTHARE PATENTION USED ID NOS: 11

SOUTHARE PATENTION USED ID NOS: 11
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
FILING DATE: 07-JUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-JUN-1995
ATMINGST/AGENT INFORMATION:
APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-JUN-1995
ATMINGST/AGENT INFORMATION:
APPLICATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944

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, ORGANISM: Homo sapiens
US-10-330-872-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   , MOLECULE TYPE: RNA
US-09-057-351-42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 2
LENGTH: 22
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4.7%; Score 21; DB 1; Length 21; 100.0%; Pred. No. 74;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Funk, Walter
APPLICANT: Funk, Walter
APPLICANT: Greider, Carol
APPLICANT: Greider, Carol
APPLICANT: Willeponteau, Bryant
ITILE OF INVENTION: RNA COMPONENT OF TELOMERASE
NUMBER OF SEQUENCES: 45
CORRESPONDENCE ADDRESS:
ADDRESSE: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: Lexington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: TELOPPY disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: Patentin Release #1.0, Version #1.30
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/08/485,778
FILING DATE: 07-21-195
CLASSIFICATION NUMBER: US 08/387,524
FILING DATE: 13-FEB-195
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-0CT-1994
PRIOR DATE: 27-0CT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 13-FEB-195
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
                PELDING DATE: 05-UN-1996
CLASSIFCATION NUMBER: US/08/660,678A
FILING DATE: 05-UN-1996
CLASSIFCATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-UL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
RESISTRATION NUMBER: 32,944
RECISTRATION NUMBER: 32,944
RECISTRATION NUMBER: 32,944
RECISTRATION INFORMATION:
TELEPHONE: (415) 576-0300
INFORMATION FFOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
LENGTH: 121 SARE DATE
LENGTH: 121 SARE DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Pred. ...
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Andrews, William H.
APPLICANT: Avilion, Ariel Athena
APPLICANT: Feng, Junil
APPLICANT: Funk, Walter
APPLICANT: Greider Carol
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 33, Application US/08485778
; Patent No. 5876979
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Best Local Similarity 100.0
Matches 21; Conservative
CURRENT APPLICATION DATA: APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: 11:
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             \mathbf{n}
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US-08-485-778-33/c
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Assays for the RNA Component of Human Telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4.7%; Score 21; DB 1; Length 21;
100.0%; Pred. No. 74;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                              COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENT PRESSENT PAPLICATION DATA:
APPLICATION NUMBER: US/08/482,115B
FILING DATE: 07-JUN-1995
FILING DATE: 07-JUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
ATPLING DATE: 27-OCT-1994
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GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Juni;

APPLICANT: Feng, Junii

APPLICANT: Feng, Junii

APPLICANT: For Mammalian H.

TITLE OF INVENTION: Mammalian Telomerase

NUMBER OF SEQUENCES: 30

CORRESPONDENCE ADRESS:

ADDRESSEE: Towneend and Townsend and Crew LLP

STREET: Two Embarcadero Center, Eighth Floor

CITY: San Francisco

CITY: San Francisco

CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patentin Release #1.0, Version #1.30
                                                         NUMBER OF SEQUENCES: 40
CORRESPONDENCE ADDRESS:
TOADRESSE: Townsend and Townsend and Crew LLi
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               184 TGCTGGCCCGTTCGCCCCTCC 204
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21 TGCTGGCCCGTTCGCCCCTCC 1
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Matches 21; Conservative
                                                                                                                                                STREET: Two Embarcade CITY: San Francisco STATE: California COUNTRY: USA
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 111
US-08-660-678A-25/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-482-115B-25
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STATE: CE
COUNTRY:
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4.7%; Score 21; DB 1; Length 21; 100.0%; Pred. No. 74; 0; Indels cive 0; Mismatches 0; Indels

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184 TGCTGGCCGGTTCGCCCTCC 204
           Query Match
Best Local Similarity 100.C
Matches 21, Conservative
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US-08-520-550A-33
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: USA

ZIP: 9411-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IB PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US 08/472,802C
FILING DATE: 07-JUN-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 07-JUL-1994
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 37-OCT-1994
ATTORNEY/AGENT INFORMATION:
NAME: SAMITCH WILLIAM M.
REFERENCE/DOCKET NUMBER: 15389-000820
TELECOMMUTICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPAX: (415) 576-0300
INPORMATION TOR SEQ ID NO: 26:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                         FILING DATE: 07-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Granahan, Patricia
REGISTRATION NUMBER: 32,227
REFERENCE/DOCKET NUMBER: CSHL94-05A4
TELECOMMUNICATION INFORMATION:
TELEPAK: 617-861-9540
INFORMATION FOR SEQ ID NO: 33:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                     184 TGCTGGCCCGTTCGCCCCTCC 204
                                                                                                                                                                                                                      ; TOPOLOGY: linear
US-08-485-778-33
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                                                                  RESULT 114
US-08-520-550A-33/C
| Sequence 3.3 Application US/08520550A
| Patent No. 601346E
| GENERAL INFORMATION:
| APPLICANT: Andrews, William H. APPLICANT: Peng, Junia
| APPLICANT: Feng, Junia
| APPLICANT: Feng, Marler
| APPLICANT: Funk, Walter
| APPLICANT: Walneda, Maria A. B. APPLICANT: Villeponteau, Bryant
| TITLE OF INVENTION: RNA Component of Telomerase
| WUMBER OF SEQUENCES: 47
| CORRESPONDENCE ADDRESS: Hamilton, Brook, Smith & Reynolds, P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4.7%; Score 21; DB 1; Length 21; 100.0%; Pred. No. 74; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/520,550A
FILING DATE: 29-AUG-1995
CLASSIPICATION: 435
PRIOR APPLICATION NUMBER: US/08/387,524
PILING APPLICATION NUMBER: US/08/387,524
PILING APPLICATION NUMBER: US/08/330,123
FILING APPLICATION NUMBER: US/08/330,123
FILING APPLICATION NUMBER: US/08/330,123
FILING APPLICATION NUMBER: US/08/330,123
FILING APPLICATION NUMBER: US/08/330,123
RIGHT INFORMATION:
NAME: Granahan, Patricia
REGISTRATION NUMBER: 32,227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     184 TGCTGGCCCGTTCGCCCTCC 204
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TELEPHONE: 617-861-6240
TELEFAX: 617-861-9540
INFORMATION FOR SEQ ID NO: 33:
SEQUENCE CHARACTERISTICS:
TENGTH: 21 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Two Militia Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: US
ZIP: 02173
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy d:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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STRANDEDNESS: single
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CITY: Lexington
STATE: MA
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RESULT 115

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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4.7%; Score 21; DB 1; Length 21; 00.0%; Pred. No. 74;
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                                              COUNTRY: USA
ZIP: 9411-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/060,523
FILING DATE: 14-APR-1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/660,678
FILING DATE: 05-JUN-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US 08/322,102
FILING DATE: 07-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: SLOYELIA JOHN R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000813US
TELEPHONE: (415) 576-0300
TELEPHONE: (415) 576-0300
TELEPHONE (415) 576-0300
TELEPHONE (HARACTERISTICS:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
SEGUENTE: TUNDER: TO BE TO THE TOWNEY.
TELEPHONE: 121 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Pred. No. 74; tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: VILLEPONTEAU, Bryant
FENG, Junli
FUNK, WALLER
ANDREWS, William H.
TITLE OF INVENTION: HUMAN TELOMERASE
NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/580,517
FILING DATE: 25-May-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         184 TGCTGGCCCGTTCGCCCTCC 204
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 100.
Matches 21; Conservative
San Francisco
California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         , MOLECULE TYPE: DNA
US-09-060-523-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-580-517-25/c
                                           COUNTRY:
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100.0%; Pred. No. 74;
tive 0; Mismatches 0; Indels
                                                                                     APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Juni
APPLICANT: Funk, Walter
APPLICANT: Funk, Walter
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                     ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/998,443
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    E: Townsend and Townsend and Crew LLP Two Embarcadero Center, Eighth Floor
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CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/660,678
FILING DATE: 05-JUN-1996
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: SLOTE11a, John R.
REFERENCE/DOCKET NUMBER: 32,944
FERENCE/DOCKET NUMBER: 32,944
FERENCE/DOCKET NUMBER: 015389-000811US
TELEPHONE: (415) 576-0300
TELEPHONE: (415) 576-0300
TELEPHONE: (415) 576-0300
INPORMATION FOR EGG ID NO: 25:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 25, Application US/09060523
Sequence 25, Application US/09060523
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junii
APPLICANT: Funk, Walter
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               184 TGCTGGCCCGTTCGCCCTCC 204
US-08-998-443-25/c; Sequence 25, Application US/08998443; Patent No. 6054575; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              / MOLECULE TYPE: DNA
US-08-998-443-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
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US-09-060-523-25/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET:
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OTHER INFORMATION: /note= "N = 5' biotinylated cytidine"
                                                                                                                                                                                                                                                                                                     4.7%; Score 21; DB 1; Length 21; 100.0%; Pred. No. 74; cive 0; Mismatches 0; Indels
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CONDITY: 1834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/833,377
FILING DATE: 04-APR-1997
CLASSIFICATION: ADATA:
APPLICATION NUMBER: US 08/510,736
FILING DATE: 04-AUG-1995
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REBERENCE/DOCKET NUMBER: 015389-001110US
TELEFHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LANDERS DATE
MANDE: ADARDED DATE
SEQUENCE CHARACTERISTICS:
LANDERS DATE
MANDE: ADARDED DATE
MANDE: ADARDED DATE
SEQUENCE CHARACTERISTICS:
LANDERS DATE
MANDE: ADARDED DATE
MANDE: ADARDED
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Patent No. 5968506

GENERAL INFORMATION:
APPLICANT: Weinrich, Scott L.
APPLICANT: Minson III, Edward M.
APPLICANT: Lichtsteiner, Serge P.
APPLICANT: Pruzan, Ronald A.
APPLICANT: Kealey, James T.
TITLE OF INVENTION: Purified Telomerase
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: -
LOCATION: 1..22
OTHER INFORMATION: /note= "oligonucleotide P3"
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 21 base pairs TYPE: mucleic acid STRANBENESS: single TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
NAME/KEY: modified_base
                                                                                                                                                                                                                                                                                                                                    Best Local Similarity 100.
Matches 21; Conservative
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linea
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                                                                                                                                                                                                                                        US-09-057-351-25
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                                                                                                                                                                                                                                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Floppy disk
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: DATE: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-JUN-1995
ATTORIEY/AGRINI INFORMATION:
ANAMERICATION NUMBER: US 08/472,802
FILING DATE: 07-JUN-1995
ATTORIEY/AGRINI INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 118
US-09-057-151-25/C
; Sequence 25, Application US/09057351
; Patent No. 6548298
; GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Unili
; APPLICANT: Funk, Walter
; TITLE OF INVENTION: Mammalian Telomerase
; TITLE OF ENGURNES: 42
; CORRESPONDENCE ADDRESS:
; ADDRESSE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STRATE: California
ZIP: 0.11
                    PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/330,123
FILING DATE: «Unknown:
ATTORNEY/AGENT INFORMATION:
NAME: Smith, william M
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 15389-000810
TELECOMMUNICATION INFORMATION:
TELEPAX: (415) 326-2400
TELEFAX: (415) 326-2422
INFORMATION FOR SEQ ID NO: 25:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000821US
FELECOMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear

MOLECULE TYPE: DNA

SEQUENCE DESCRIPTION: SEQ ID NO: 25:
US-09-580-517-25
   CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           184 TGCTGGCCCGTTCGCCCCTCC 204
                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity
Matches 21; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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RESULT 122
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                                        DB 1; Length 22; 78;
                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:

MEDIUM TYPE: 5109Py disk

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

COMPUTER: APELICATION DATA:

APPLICATION NUMBER: US/08/330,123A

FILING DATE: 27-0CT-1994

FILING DATE: 107-0CT-1994

ATTORNEY/AGENT INFORMATION:

NAME: SMICH MILLIAM M

REGISTRATION NUMBER: 30,223

REFERENCE/DOCKET NUMBER: 15389-000810

TELECOMMUNICATION INFORMATION:

MATERIALICATION INFORMATION:

NAME: ATTORNEY/AGENT NUMBER: 30,223

REFERENCE/DOCKET NUMBER: 15389-000810
                                                                                                                                                                                                                                                                                                                        APPLICANT: VILLEBONTEAU, Bryant
APPLICANT: FENG, Junii
APPLICANT: FENG, Junii
APPLICANT: FUNK, Walter
TITLE OF INVENTION: HUMAN TELOMERASE
NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend Khourie and Crew
                                      4.7%; Score 21; DB 100.0%; Pred. No. 78; tive 0; Mismatches
                                                                                                                                                                                                                                                               Sequence 4, Application US/08330123A
Patent No. 5583016
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-330-123A-7/c
; Sequence 7, Application US/08330123A
; Patent No. 5583016
; GENERAL INFORMATION:
APPLICANT: VILLEPONTEAU, Bryant
APPLICANT: FENG, Junil
                                                                                                                        46 CTAACCCTAACTGAGAAGGGC 66
                                                                                                                                               22 CTAACCCTAACTGAGAAGGGC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          41 TTTGTCTAACCCTAACTGAG 60
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TELEFAX: (415) 326-2422
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              379 Lytton Avenue
                                    Query Match
Best Local Similarity 100.0
Matches 21; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Palo Alto
STATE: California
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear / MOLECULE TYPE: RNA US-08-330-123A-4
                                                                                                                                                                                                                                             US-08-330-123A-4/c
US-08-833-377-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET:
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APPLICANT: Feng, Junia
APPLICANT: Feng, Junia
APPLICANT: Feng, Junia
APPLICANT: Andrews William H.
TITLE OF INVENTION: Assays for the RNA Component of Human
TITLE OF INVENTION: Telomerase
NUMBER OF SEQUENCES: 40
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:

MEDIUM TYPE: Floppdisk

MEDIUM TYPE: Floppdisk

COMPUTER: IBM PC compatible

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/330,123A

FILING DATE: 27-CCT-1994

FLING DATE: 17-CCT-1994

ATTORNEY/AGENT INFORMATION:

NAME: SMIGHT WILLIAM M

REGISTRATION NUMBER: 30,223

REFERENCE/DOCKET NUMBER: 15389-000810

TELECOMMUNICATION INFORMATION:

TOTAL COMPUTER: TOTAL TOTA
APPLICANT: FUNK, Walter
APPLICANT: ANDREWS, William H.
TITLE OF INVENTION: HUMAN TELOMERASE
NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend Khourie and Crew
STREET: 379 Lytton Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPDY disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4.4%; Score 20; DB 100.0%; Pred. No. 83; ive 0; Mismatches
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Patent No. 5776679
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 GGTTGCGGAGGGTGGGCCTG 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (415) 326-2400
TELEPAX: (415) 326-2422
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 100.0
Matches 20; Conservative
                                                                                                                                                                                                                                                                 STREET: 379 Lytton
CITY: Palo Alto
STATE: California
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-482-115B-7/c
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Gaps
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83;
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                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
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ZIP: 94111-3834

ZIP: 94111-3834

COMPUTER READABLE FORM:

MEDIUW TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PATCHION DATA:

APPLICATION NUMBER: US/08/660,678A

FILING DATE: 05-JUN-1996

CLASSIFICATION 1435

PRIOR APPLICATION NUMBER: US/08/330,123

FILING DATE: 27-OCT-1994

PRIOR APPLICATION NUMBER: US/08/330,123

FILING DATE: 27-OCT-1994

PRIOR APPLICATION NUMBER: US/08/330,123

FILING DATE: 27-OCT-1994

ATTONNEY/AGENT INFORMATION:

ANAME: ACCOUNT ACCOUNT ACCOUNTS ANAME: ACCOUNTS ACCOUN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Townsend and Townsend and Crew LLP
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                                                                                                                                                                                                                                                                                                                                              Query Match

4.4%; Score 20; DB
Best Local Similarity 100.0%; Pred. No. 83;
Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   015389-000811US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junil
APPLICANT: Feng, Junil
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ANDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-660-678A-26/c; Sequence 26, Application US/08660678A; Patent No. 5837857.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 GGTTGCGGAGGGTGGGCCTG 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     41 TTTGTCTAACCCTAACTGAG 60
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REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 01
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 26:
SEQUENCE CHARACTERISTICS:
       (415) 576-0300
                                   INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          San Francisco
: California
                                                                                                                                                                                                                 linear
                                                                                                                                                                                                                                              MOLECULE TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE:
                                                                                                                                                                                                              TOPOLOGY:
                                                                                                                                                                                                                                                                                 US-08-660-678A-7
       TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET:
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Pred. No. 83;
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/660,678A
FILING DATE: 05-JUN-1996
CLASSIFICATION: 435
PRIOR APPLICATION NUMBER: US/08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
ATONINY/AGENT INFORMATION:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
ATONINY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/POCKET NUMBER: 32,944
REGISCOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
PROFILE APPLICATION NUMBER: US 08/330,123
APPLICATION NUMBER: US 08/330,123
ATORNEY AGENT: 1NFORMATION:
NAME: STOOTHIA, JOHN R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000830US
TELEPAN: (415) 576-0200
TELEPAN: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US/08/482,115B
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 7, Application US/08660678A
Patent No. 5837857
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GGTTGCGGAGGGTGGGCCTG 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GGTTGCGGAGGGTGGGCCTG 1
APPLICATION NUMBER: US/01
FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            , MOLECULE TYPE: DNA
US-08-482-115B-7
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ZIP: 94111-3834
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 123
US-08-660-678A-7/c
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Gaps
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83;
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Sequence 6, Application US/08770565
; Sequence 6, Application US/08770565
; Patent No. 5846727
; Patent No. 5846727
; APPLICANT: Kim, Nam Woo
; APPLICANT: Kaley, James T.
APPLICANT: Realey, James T.
APPLICANT: Realey, James T.
APPLICANT: Meinrich, Scott L.
; TILLE OF INVENTION: Methods for Detecting the RNA TITLE OF INVENTION: Telomerase
NUMBER OF SEQUENCES: 26
CORRESPONDENCES 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: TOWNSEND and TOWNSEND and CREW LLP STREET: Two Embarcadero Center, 8th Floor CITY: San Francisco
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/770,565 FILING DATE: 20-DEC-1996 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4.4%; Score 20; DB 100.0%; Pred. No. 83; tive 0; Mismatches
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           361 AGGCCGCAGGAAGAACG 380
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CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-770-565-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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Patent No. 5846723
GENERAL INFORMATION:
APPLICANT: Kim, Nam Woo
APPLICANT: Kealey, James T.
APPLICANT: Realey, James T.
APPLICANT: Weinrich, Scott L.
TITLE OF INVENTION: Methods for Detecting the RNA Component of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 20; DB 1; Length 20;
Pred. No. 83;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                 Sequence 29, Application US/08660678A

Sequence 29, Application US/08660678A

GENERAL INFORMATION:

APPLICANT: Villeponteau, Bryant
APPLICANT: Peng, Junli

APPLICANT: Andrews, William H.

TITLE OF INVENTION: Mammalian Telomerase

NUMBER OF SEQUENCES: 30

CORRESPONDENCE ADDRESS:

ADDRESSEE: Townsend and Townsend and Crew LLP

STREET: Two Embarcadero Center, Eighth Floor

CITY: San Francisco

SCATAE: California
                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PAPELICATION NUMBER: US/08/660,678A
FILING DATE: 05-JUN-1996
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: SLOCATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000811US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4.4%; bcc...
100.0%; pred. No. c...
... 0; Mismatches
             20 TIGITIAACCTAACTGAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              , MOLECULE TYPE: RNA
US-08-660-678A-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               linear
                                                                                      RESULT 125
US-08-660-678A-29/c
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Gaps
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APPLICANT: Kim, Nam Woo
APPLICANT: Wi, Fred
APPLICANT: World
APPLICANT: Weinrich, Scott L
APPLICANT: Pruzan, Ronald
APPLICANT: Pruzan, Scott L.
TITLE OF INVENTION: Telomerase
NUMBER OF SEQUENCES: 26
NUMBER OF SEQUENCES: 26
STREET: Two Embarcadero Center, 8th Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: CALLYLLINGS

COMPUTRY: USA

ZIP: 9411-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN DATA:
CURREL PATENTIN DATA: 20-DEC-1996
CLARSIFICATION NUMBER: US/08/770,565
FILING DATE: 20-DEC-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERRACE/DOCKET NUMBER: 015389-002300US
TELEPAX: 415-576-0300
TELEFAX: 415-576-0300
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
CWATH. 20 base pairs
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APPLICANT: Avilion, Ariel Athena
APPLICANT: Feng, Junil
APPLICANT: Feng, Junil
APPLICANT: Greider, Carol
APPLICANT: Marhuenda, Maria Antonia Blasco
APPLICANT: Wileponteau, Bryant
TITLE OF INVENTION: RNA COMPONENT OF TELOMERASE
NUMBER OF SEQUENCES: 45
290 CTGCCACCGCGAAGAGTTGG 309
                                                                                                                                                 ; Sequence 13, Application US/08770565; Patent No. 5846723
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     159 TCTAGAGCAAACAAAAATG 178
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                             20 CTGCCACCGCGAAGAGTTGG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; TOPOLOGY: linear; MOLECULE TYPE: DNA US-08-770-565-13
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nes 20; Conserve
                                                                                                      RESULT 129
US-08-770-565-13/c
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                                                                                                                                                                                                                                                                                                                                                                     Gaps
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Fatent No. 5846723
GENERAL INFORMATION:
APPLICANT: Kin, Nam Woo
APPLICANT: Wein, Fred
APPLICANT: Weinzen, Scott L.
TITLE OF INVENTION: Methods for Detecting the RNA Component of
TITLE OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSE: TWO SEMBARCAGERO CENTER, 8th Floor
STREET: Two Embarcadero Center, 8th Floor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE PATENTIN Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/770,565 FILING DATE: 20-DEC-1996
CLASSIFICATION: 435
ATTONREY/AGENT INPORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REBRENCE/POCKET NUMBER: 015389-002300US
TELECOMMULCATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: Two Embarcadero Center, 8th Floor
CITY: San Francisco
STATE: California
           32,944
RR: 015389-002300US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; Pred. ...
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                           300 GAAGAGTIGGGCTCTGTCAG 319
        REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 0153(
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-576-0200
TELEPKX: 415-576-0300
INFORMATION FOR SEQ ID NO: 6: SEQUENCE CHARACTERISTICS:
LENGINE CHARACTERISTICS:
LENGINE CHARACTERISTICS:
TYPE: Nucleic acid
STRANDENNES: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                    20 GAAGAGTIGGGCTCTGTCAG 1
                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 100.
Matches 20; Conservative
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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Matches 20; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                    ; TOPOLOGY: linear; MOLECULE TYPE: DNA US-08-770-565-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             linear
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                 CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/485,778
FILING DATE: 07-JE-1995
CLASSIFICATION NUMBER: US/08/485,778
FILING DATE: 13-FEB-1995
PRIOR APPLICATION NUMBER: US 08/387,524
FILING DATE: 13-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
ATTONENTY/AGENT INFORMATION:
NAME: Granahan, Patricia
REGISTRATION NUMBER: 32,227
REFRENCE/DOCKET NUMBER: 32,227
REFRENCE/DOCKET NUMBER: 32,227
TELECOMMUNICATION INFORMATION:
TELEBRAK: 617-861-6240
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/472,802C
FILING DATE: 07-JUN-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 8, Application US/08472802C
Fatent No. 5958680
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junil
APPLICANT: Andrews, William H.
ITLE OF INVENTION: Mammalian Telomerase
ITLE OF SEQUENCES: 44
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew ilp
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
CITY: San Francisco
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC_DOS/MS-DOS
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ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 9
SEQUENCE CHARACTERISTICS:
TRUGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; TOPOLOGY: linear
US-08-485-778-9
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ZIP: 94111-3834
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 132
US-08-472-802C-8/c
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COUNTRY: US
STATE: WA
COUNTRY: US
CITY. US
CITY: US
CITY
CITY: US
CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: Two Militia Drive
CITY: Lexington
                                                                                                                                                                                                                COMPUTER: TEADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: TEM PC compatible
CORPERATION SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/485,778
FILING DATE: 07-JE-1995
CLASSIFFACATION: 435
PRIOR APPLICATION ADATA:
APPLICATION NUMBER: US 08/387,524
FILING DATE: 13-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-CCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
REGISTRATION NUMBER: 32,227
REFERENCE/DOCKET NUMBER: CSHL94-05A4
TELECOMMUNICATION INFORMATION:
TELEBRHONE: 617-861-6540
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CORRESPONDENCE ADDRESS:
ADDRESSE: Hamilton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Andrews, William H.
Avilion, Ariel Athena
Feng, Junli
Funk, Walter
Greider, Carol
Marhuenda, Maria Antonia Blasco
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Patent No. 5876979
GENERAL INFORMATION:
APPLICANT: Andrews, William H.
APPLICANT: Feng, Junli
APPLICANT: Feng, Junli
APPLICANT: Feng, Junli
APPLICANT: Greider, Carol
APPLICANT: Greider, Carol
APPLICANT: Mathuenda, Maria Antoni
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 617-861-9540
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 100.
Matches 20; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
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02173
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02173
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US-08-485-778-6
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                                                                                          CITY: Lex
STATE: MA
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ZIP: 94111-3834

ZIP: 94111-3834

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

CORPUTER: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/472,802C

FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 133
US-08-472-802C-41/C
; Sequence 41, Application US/08472802C
; Patent No. 5958680
; GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Unnli am H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 44
NUMBER OF SEQUENCES: 44
NUMBER OF SEQUENCES: 44
STREET: Two Embarcadero Center, Eighth Floor
STREET: Two Embarcadero Center, Eighth Floor
        REFERENCE/DOCKET NUMBER: 15389-000820
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDENNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 07-UN-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-UL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
REFENCE/DOCKET NUMBER: 15389-00082
TELECOMMUNICATION INPORMATION:
TELECOMMUNICATION INPORMATION:
                                                                                                                                                                                                                                                                                                                                                                                      2 GGTTGCGGAGGGTGGGCCTG 21
REGISTRATION NUMBER: 30,223
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INFORMATION FOR SEQ ID NO: 41:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                TOPOLOGY: linear MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                        Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: Two concerts CITY: San Francisco cTATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; MOLECULE TYPE: RNA
US-08-472-802C-41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             linear
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4.4%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 83;

Query Match Best Local Similarity

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Gaps
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Indels
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CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/833,377
FILING DATE: 04-APR-1997
CLASSIFICATION: 435
PRIOR APPLICATION NUMBER: US/08/510,736
FILING DATE: 04-ADR-1995
ATFORNEY/AGENT INPORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REGISTRATION STORET NUMBER: 32,944
REGISTRATION STORET NUMBER: 32,944
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REGISTRATION STORET STORET STORET STORE STO
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                                                                                                                                                                                                                                                                                                                                  Sequence 7, Application US/08833377
Patent No. 5968506
GENERAL INFORMATION:
APPLICANT: Meinrich, Scott L.
APPLICANT: Atkinson III, Edward M.
APPLICANT: Lichtsteiner, Serge P.
APPLICANT: Pruzan, Ronald A.
APPLICANT: Pruzan, Ronald A.
APPLICANT: Realey, James T.
TITLE OF INVENTION: Purified Telomerase
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
0; Mismatches
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LOCATION: 1..20
OTHER INFORMATION: /note= "Oligo 14ab"
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            361 AGGCCGCAGGAAGAGGAACG 380
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20; Conservative
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ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nucleic acid
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US-08-520-550A-6/c
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Matches
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DB 1; Length 20;
83;
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                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: RADABLE FORM:
MEDIUM TYPE: RADABULE FORM:
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/520,550A
FILING DATE: 29-AUG-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/387,524
FILING DATE: 13-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FRIDR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FRIING DATE: 27-OCT-1994
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: GRANAHAIN PARTICIA
REGISTRATION NUMBER: 32,227
REFERENCE/POCKET NUMBER: 32,227
REFERENCE/POCKET NUMBER: 32,227
REFERENCE/POCKET NUMBER: GSHL94-05A3B
TELEFRAX: 617-861-6240
TANDAMATICATION NUMBER: OF 17-861-6240
TELEFRAX: 617-861-6240
  Brook, Smith & Reynolds, P.C.
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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TELEFAX: 617-861-9540
INFORMATION FOR SEQ ID NO: 9: SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
3: Hamilton, Broo.
Two Militia Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
          STREET: Two ...
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US-08-520-550A-9
  ADDRESSEE:
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                                                                                  COUNTRY:
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                                                   APPLICANT: Funk, Walter
APPLICANT: Greider, Carol
APPLICANT: Marhuenda, Maria A. B.
APPLICANT: Marhuenda, Maria A. B.
TITLE OF INVENTION: RNA Component of Telomerase
NUMBER OF SEQUENCES: 47
CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: Two Militia Drive
CITY: Lexington
                                                                                                                                                                                                                                                                                     COMPTRY: US

CONTRY: US

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IS FORM:
MEDIUM TYPE:
COMPATING SYSTEM: PC_DOS/MS_DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/520,550A
FILING DATE: 29-AUG-1995
CLASSIPICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/387,524
FILING DATE: 13-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-CT-1994
FILING DATE: 27-CT-1994
ATTORNEY/AGENT INFORMATION:
NAME: Grandhan, Patricia
REGISTRATION NUMBER: 32,227
REFERENCE/DOCKET NUMBER: 32,227
REFERENCE/DOCKET NUMBER: CSHL94-05A3B
TELEPHONE: 617-661-6640
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Andrews, William H.
APPLICANT: Andrews, William H.
APPLICANT: Peng, Junii
APPLICANT: Feng, Junii
APPLICANT: Greider, Carol
APPLICANT: Marhuenda, Maria A. B.
APPLICANT: Vileponteau, Bryant
TITLE OF INVINION: RNA COMPONENT OF SEQUENCES: 47
CORRESPONDENCE ADDRESS:
                                     Feng, Junli
Funk, Walter
Greider, Carol
Marhuenda, Maria A. B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-08-520-550A-9/c
; Sequence 9, Application US/08520550A
; Patent No. 6013468
; GENERAL INFORMATION:
; APPLICANT: Andrews, William H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         41 TITGICTAACCCTAACTGAG 60
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  Andrews, William H. Avilion, Ariel A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: 6: SEQUENCE CHARACTERISTICS: LENGTH: 20 base pairs TYPE: nucleic acid STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity 100.
Matches 20; Conservative
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                                                                                                                                                                                                                                   CITY: Lexin
STATE: MA
COUNTRY: U
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other nucleic acid (PNA), desc = "peptide nucleic acid (PNA), where (deoxy/tibose-phosphate linkages are replaced by N-(2-aminoethyl)glycine units linked to nucleotide bases via glycine amino N-trough a methylenecarbonyl linker"
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                                                                                                                                                                                                                                                                                                                                                                                                     4.4%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 83; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Juuli
APPLICANT: Funk, Walter
APPLICANT: Funk, Walter
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 30
CORRESSONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/998,443 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/660,678
FILING DATE: 05-UN-1996
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-0CT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-UL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTATION NUMBER: 32,944
REGISTATION NUMBER: 015389-000811US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4.4%; Score 20;
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ZIP: 9411-3834
COMPUTER READABLE FORM:
COMPUTER: Floppy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          46 CTAACCCTAACTGAGAAGGG 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-998-443-7/c
; Sequence 7, Application US/08998443
; Patent No. 6054575
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TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLLGY: linear
MOLECULE TYPE: other nucl
DESCRIPTION: where (deo
DESCRIPTION: where (deo
DESCRIPTION: 00 pairs
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Best Local Similarity 100.0
Matches 20; Conservative
SEQUENCE CHARACTERISTICS:
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STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                               US-08-838-545-41
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /desc = "peptide nucleic acid (PNA), where (deoxy)ribose-phosphate linkages are replaced by N-(2-aminoethyl)glycine units linked to nucleotide bases via Blycine amino nitrogen through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 20;
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SITALE: CALIFOLITIA

ZIP: 94111-3834

COMPUTER READABLE FORM:
MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN RE-BESE #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
CLASSIFICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
APPLICATION NUMBER: US/08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: SLOCATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0200
INFORMATION FOR SEQ ID NO: 41:
                  ATTORNAY OF THE TOTAL OF THE TO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
4.4%; Score 20; DB 1;
Best Local Similarity 100.0%; Pred. No. 83;
Matches 20; Conservative 0; Mismatches
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Patent No. 6046307
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       CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Shay,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: 1in
MOLECULE TYPE:
DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 138
US-08-838-545-41/c
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DB 1; Length 20;
83;
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                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/998,443 FILING DATE:
                                                   APPLICANT: Funk, Walter
APPLICANT: Funk, Walter
APPLICANT: Andrews, William H.
ITILE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSEE: Twomsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Townsend and Townsend and Crew LLP STREEF: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
PRIOR APPLICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/660,678
FILING DATE: 05-UN-1996
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-CCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-UL-1994
ATTORNEY/AGENT INFORMATION:
NAME: SECRETIAL JOHN R.
REGISTRATION NUMBER: 32,944
RECENENCE/DOCKET NUMBER: 32,944
RECENENCE/DOCKET NUMBER: 015389-000811US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0200
INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 Asse pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      | Sequence 7, Application US/09060523 | Patent No. 6258535 | GENERAL INPORMATION: APPLICANT: Ford, Juni APPLICANT: Feng, Juni APPLICANT: Feng, Juni APPLICANT: From, Walter APPLICANT: Ford, Walliam H. TITLE OF INVENTION: Mammalian Telomerage NUMBER OF SEQUENCES: 25 | CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 4.4%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 83; Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                       PC-DOS/MS-DOS
         Villeponteau, Bryant
Feng, Junli
Funk, Walter
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                                                                                                                                                                                                                                                             COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: INP PC COMPATIBLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                     OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TOPOLOGY: linear; MOLECULE TYPE: RNA
US-08-998-443-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-060-523-7/c
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                                    Gaps
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Pred. No. 83;
                                    0; Indels
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                                                                                                                                                                                                                                  Sequence 26, Application US/08998443
; Sequence 26, Application US/08998443
; Patent No. 6054575
; GENERAL INFORMATION:
    APPLICANT: Feng, Juni,
    APPLICANT: Feng, Juni,
    APPLICANT: Feng, Juni,
    APPLICANT: Form, Walter
    APPLICANT: Andrews, William H.
    TITLE OF INVENTION: Mammalian Telomerase
    NUMBER OF SEQUENCES: 30
    CORRESPONDENCE ADDRESS:
    ADDRESSER: Townsend and Townsend and Crew LLP
    STREET: Two Embarcadero Center, Eighth Floor
    CITY: San Francisco
    STATE: Canifornia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
ZIP: 94111-3834
COMPUTER REDABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATCHIT Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/998,443
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRILING DATE:

CLASSIFICATION
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/660,678
FILING DATE: 05-JUN-1996
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
ATORNEY AGENT INFORMATION:
NAME: SCOFELLA, JOHN R.
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
ATORNEY AGENT INFORMATION:
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 26:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 Asserpairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
4.4%; Score 20; UB
Best Local Similarity 100.0%; Pred. No. 83;
100.0%; Pred. ...
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; Sequence 29, Application US/08998443
; Patent No. 6054575
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           41 TTTGTCTAACCCTAACTGAG 60
                                                                                2 GGTTGCGGAGGGTGGGCCTG 21
                                                                                                                            20 dárraccadadaraccici 1
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         Best Local Similarity 100. Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     , MOLECULE TYPE: DNA
US-08-998-443-26
                                                                                                                                                                                                   RESULT 140
US-08-998-443-26/c
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Gaps ö

94111-3834

COUNTRY:

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other nucleic acid (PNA),

/desc = "peptide nucleic acid (PNA),
where (deoxy(ribose-phosphate linkages are replaced by
N-(2-aminoethy)ajvaine units linked to nucleotide bases via
glycine amino N through a methylenecarbonyl linker"
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MEDIUM TYBES FLORDY disk
COMPUTER READABLE FORM:

MEDIUM TYBES FLORDY disk
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC COMPATIBLE FORM:
COMPUTER: DATE: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/09/580,517
FILING DATE: 25-May-2000
CLASSIFICATION NUMBER: 08/330,123
APPLICATION NUMBER: 08/330,123
APPLICATION NUMBER: 08/330,123
REFERENCE/DOCKET NUMBER: 30,233
REFERENCE/DOCKET NUMBER: 15389-000810
TELECOMMUNICATION INPORMATION:
MATE SMITCH 15, 256.2300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
ADDRESSER: Townsend and Townsend Khourie and Crew
STREET: 379 Lytton Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
4.4%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 83;
Matches 20; Conservative 0; Mismatches 0; Indels
     APPLICATION NUMBER: US 08/838,545
FILING DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: SCOTELLA, UDAN R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 41:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-580-517-4/c
; Sequence 4, Application US/09580517
; Patent No. 632039
; GENERAL INFORMATION:
; APPLICANT: VILLEPONTEAU, Bryant FRNG, Juni;
; FUNK, Walter FUNK, Walter FUNK, Walter FUNK, Walter FUNK, Walter FUNK WILLER OF INVENTION: HUMAN TELOMERASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (415) 326-2400
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              46 CTAACCCTAACTGAGAAGGG 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20 CTAACCCTAACTGAGAAGGG 1
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STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
MOLECULE TYPE: othe
DESCRIPTION: /des
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
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US-09-349-532-41/c

is Sequence 41, Application US/09349532

is Patent No. 6294650

is GENERAL INFORMATION:

is APPLICANT: Shay, Jerry W.

APPLICANT: Wright, Woodring E.

APPLICANT: Piatysek, Mieczyslaw A.

APPLICANT: Orey, David R.

APPLICANT: No. 629450ton, James C.

ITILE OF INVENTION: Peptide Nucleic Acids

ITILE OF INVENTION: Peptide Nucleic Acids

NUMBER OF SEQUENCES: 60

CORRESPONDENCES: 60

CORRESPONDENCES: 1 Two Embarcadero Center, Eighth Floor

CITY: San Francisco

STATE: California
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                                                                                  CURRENT APPLICATION DATA:

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/060,523

FILING DATE: 14-APR-1998

CLASSIPICATION: 536

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/08/660,678

FILING DATE: 05-UN-1996

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/330,123

FILING DATE: 27-OCT-1994

PRIOR APPLICATION NUMBER: US 08/330,123

FILING DATE: 27-OCT-1994

APPLICATION NUMBER: US 08/320,123

FILING BATE: 07-OCT-1994

APPLICATION NUMBER: US 08/320,102

FILING BATE: 07-UL-1994

ATONEY/AGENT INFORMATION:

REGISTRATION NUMBER: 32,944

REGISTRATION NUMBER: 015389-000813US

TELEPRANCE/DOCKET NUMBER: 015389-000813US

TELEPRANCE/COCKET NUMBER: 015389-000813US

TELEPRANCE/COCKET NUMBER: 015389-000813US

TELEPRANCE/COCKET NUMBER: 015389-000813US

TELEPRANCE/COCKET NUMBER: 015389-000813US

TELEPRANCE CHARACTERISTICS:

LEMOTH: 02 Dates pairs

TVVB: 12 Dates pairs
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
CURRENT APPLICATION DATA:
FILING DATE:
FLASSIFICATION:
CLASSIFICATION:
PRIOR APPLICATION DATA:
                     MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20 GGTTGCGGAGGTGGGCCTG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
WOLECULE TYPE: DNA
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Townsend and Townsend Khourie and Crew STREET: 379 Lytton Avenue CITY: Palo Alto STATE: California
                                                                                                                                      Query Match
4.4%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 83;
Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4.4%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 83; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
SEATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/330,123
FILING DATE: «Unknown-
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 15389-000810
TELECOMMUNICATION:
TELECOMMUNICATION:
TELECHONE: (415) 326-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/580,517
FILING DATE: 25-May-2000
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ANDREWS, William H.
TITLE OF INVENTION: HUMAN TELOMERASE
NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: DNA SEQUENCE DESCRIPTION: SEQ ID NO: 7: US-09-580-517-7
                                          MOLECULE TYPE: RNA
SEQUENCE DESCRIPTION: SEQ ID NO: 4:
US-09-580-517-4
                                                                                                                                                                                                                                                                                                                                      US-09-580-517-7/c
; Sequence 7, Application US/09580517
; Patent No. 6320039
; GENERAL INFORMATION:
; APPLICANT: VILLEPONTEAU, Bryant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 GGTTGCGGAGGGTGGCCTG 21
                                                                                                                                                                                                                      41 TTTGTCTAACCCTAACTGAG 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FELEFAX: (415) 326-2422
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
STRANDEDNESS: single
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                             FENG, Junli
FUNK, Walter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: 7
                                        TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
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ZIP: 94301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
Matches 20; Conserva
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RESULT 146

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Felease #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351 FILING DATE: 08-APR-1994
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                                                                                                                                                                                                                                                                                                          STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNDRER:
APPLICATION UNDRER:
FILING DATE: 07-UL-1994
PRIOR APPLICATION UNDRER: US 08/330,123
APPLICATION UNDRER: US 08/330,123
FILING DATE: 27-CT-1994
PRIOR APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-UN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTATION NUMBER: 32,944
REGISTATION NUMBER: 32,944
REGISTATION NUMBER: 32,944
REGISTATION NUMBER: 32,944
REGISTATION NUMBER: 015389-000821US
TELECOMMUNICATION INFORMATION:
US-09-057-351-7/c

Sequence 7, Application US/09057351

Patent No. 6548298

GENERAL INFORMATION:

APPLICANT: Villeponteau, Bryant

APPLICANT: Feng, Junia

APPLICANT: Funk, Walter

APPLICANT: Andrews, William H.

TITLE OF INVENTION: Mammalian Telomerase

NUMBER OF SEQUENCES: 42

CORRESPONDENCE ADDRESS:

ADDRESSEE: Townsend and Townsend and Crew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Unuli
APPLICANT: Feng, Unuli
APPLICANT: Funk, Walter
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 100.0%; Fred. No. 83;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-057-351-40/c
; Sequence 40, Application US/09057351
; Patent No. 6548298
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2 GGTTGCGGAGGGTGGGCCTG 21
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                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0200
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: DNA
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Patent No. 6046307
GENERAL INFORMATION:
APPLICANT: Wight, Woodring E.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Orey, David R.
APPLICANT: No. 6046307ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSES: Townsend and Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1; Length 19; 93;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSER: ADDRESS: ADDRESSER: ADDRESSER: ADDRESSER: Townsend and Townsend and Crew Lip STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California COUNTRY: USA ZIF: 94111-3834

COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatible COMPUTER: PIEM PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997

PRILING DATE: 09-APR-1996

ATTORNEY/AGENT INFORMATION: ADDRESSERIES OF APR-1996

ATTORNEY/AGENT INFORMATION: NUMBER: US/08/818,545

PRILING DATE: 09-APR-1996

ATTORNEY/AGENT INFORMATION: NUMBER: US/08/818,545

PRILING DATE: US/08/811996

ATTORNEY/AGENT INFORMATION: US/08/818,545
                                                                                                ATTORNEY/AGENT INFORMATION:

NAME: Storella, John R.
REGISTATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-002300US
TELEFONE: 415-576-0200
TELEFAX: 415-576-0300
TYPE: DAGE PARACTER: 511CS: LENGTH: 19 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4.2%; Scor.
100.0%; Pred. No. 20.
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
IFLEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 60:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
                           APPLICATION NUMBER: US/08/770,565
FILING DATE: 20-DEC-1996
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          148 CCACCGTTCATTCTAGAGC 166
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Best Local Similarity 100.0'
Matches 19; Conservative
  CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                            ; TOPOLOGY: linear; MOLECULE TYPE: DNA US-08-770-565-9
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STRANDEDNESS:
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APPLICANT: Kim, Nam Woo
APPLICANT: Kim, Nam Woo
APPLICANT: Wu, Fred
APPLICANT: Wu, Fred
APPLICANT: Wealey, James T.
APPLICANT: Wealey, James T.
APPLICANT: Wealnrich, Scott L.
TITLE OF INVENTION: Methods for Detecting the RNA Component of TITLE OF INVENTION: Telomerase
CORRESPONDENCE ADDRESS:
ADDRESSEE: TOWNSEND and TOWNSEND and CREW LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4.4%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 83; cive 0; Mismatches 0; Indels
                   CUNTATE: CCA...
COUNTATE: CCA...
COUNTATE: USA
ZIP: 94111-3834
COOUNTER: EADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
FILING APPLICATION: 435
PRIOR APPLICATION NUMBER: US/08/272,102
FILING APPLICATION NUMBER: US/08/330,123
FILING APPLICATION NUMBER: US/08/330,123
FILING APPLICATION NUMBER: US/08/472,802
FILING APPLICATION NUMBER: US/08/472,802
FILING APPLICATION NUMBER: US/08/472,802
FILING APPLICATION NUMBER: US/08/472,802
FILING DATE: 07-UN-1995
ATPORNEY/AGENT INFORMATION:
APPLICATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REFERRENCE/DOCKET NUMBER: 015389-000821'
TELEPAAN: (415) 576-0200
INFORMATION FOR SEQ ID NO: 40:
SEQUENCE CHARACTERISTICS:
TWGTH: 20 base pairs
TWGTH: 20 base pairs
TWGTH: 20 base pairs
TWGTH: 20 base pairs
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/WS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
Two Embarcadero Center, Eighth Floor
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CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 9, Application US/08770565
Patent No. 5846723
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              41 TITGICTAACCCTAACTGAG 60
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Best Local Similarity 100.(
Matches 20; Conservative
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Gaps

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US-09-018-125-2
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                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Shay, Jerry W.
APPLICANT: Wright, Woodring E.
APPLICANT: PietyPszek, Mieczyslaw A.
APPLICANT: Corey, David R.
APPLICANT: Corey, David R.
APPLICANT: No. 6294650ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by
TITLE OF INVENTION: Peptide Nucleic Acids
                                                                                  Query Match 4.2%; Score 19; DB 1; Length 19; Best Local Similarity 78.9%; Pred. No. 93; Matches 15; Conservative 4; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4.2%; Score 19; DB 1; Length 19; 78.9%; Pred. No. 93;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 94111-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IB P Looppy disk
COMPUTER: PROC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/838,545
FILING DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
RECISTRATION NUMBER: 32,946
RECISTRATION FOR SEQ ID NO: 60:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 Assepairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4; Mismatches
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APPLICATION NUMBER: US/09/349,532
                                                                                                                                                                                                                                                                                                                            Sequence 60, Application US/09349532
Patent No. 6294650
GENERAL INFORMATION:
APPLICANT: Wright, Woodring E.
                                                                                                                                                                                                   |:|:||||:|||:|||||
1 GUCUAACCCUAACUGAGAA 19
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; TOPOLOGY: linear
; MOLECULE TYPE: RNA (genomic)
US-08-818-545-60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: RNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15, Conservative
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CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bingle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: sing
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Best Local Similarity
Matches 15, Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: Ca
                                                                                                                                                                                                                                                                                                               US-09-349-532-60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-349-532-60
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RESULT 151

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                                                                                                               APPLICANT: Silverman, Robert H.
APPLICANT: Silverman, Robert H.
APPLICANT: Silverman, Robert H.
APPLICANT: Kondo, Seiji
APPLICANT: Cowell, John K.
APPLICANT: Cowell, John K.
APPLICANT: Torrence, Paul F.
TITLE OF INVENTION: RNASE L ACTIVATORS AND ANTISENSE OLIGONUCLEOTIDES
TITLE OF INVENTION: EFFECTIVE TO TREAT TELOMERASE-EXPRESSING MALIGNANCIES
FILE REFERENCE: 8656-022
CURRENT APPLICATION NUMBER: US/09/018,125A
CURRENT FILING DATE: 1999-02-03
EARLIER APPLICATION NUMBER: 60/044,507
EARLIER FILING DATE: 1997-04-21
NUMBER OF SEQ ID NOS: 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Jos-08-833-377-14/C

Sequence 14, Application US/08833377

Patent No. 5968506

GENERAL INFORMATION:

APPLICANT: Weinrich, Scott L.

APPLICANT: Weinrich, Scott E.

APPLICANT: Lichtsteiner, Serge P.

APPLICANT: Vasserot, Alain P.

APPLICANT: Pruzan, Ronald A.

APPLICANT: Puzan, Ronald A.

APP
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/833,377
FILING DATE: 04-APR-1997
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: 0.5389-001110US TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.2%; Score 19; DB 100.0%; Pred. No. 93; iive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 04-APR-1997
CLASSIFICATION: 435
FILOR APPLICATION DATE:
APPLICATION NUMBER: US 08/510,736
Sequence 2, Application US/09018125A Patent No. 6468983 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              76 GIGCITITGCICCCGCGC 94
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ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Ver. 2.0 SEQ ID NO 2 LENGTH: 19 TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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Matches 19; Conserv
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nucleic acid
  linear
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Best Local Similarity
                MOLECULE TYPE: O DESCRIPTION: V DESCRIPTION: V DESCRIPTION: N DESCRIPTION: DESCRIPTION: E
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DESCRIPTION:
DESCRIPTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DESCRIPTION:
DESCRIPTION:
                                                                                                                                                                                                                                                                                                                                                                     US-09-349-532-9/c
  TOPOLOGY:
                                                                                                                        US-08-838-545-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-349-532-9
                                                                                                                                                                Query Match
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Matches
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APPLICANT: Wright, Woodring E.
APPLICANT: Wright, Woodring E.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Corey, David R.
APPLICANT: No. 6046307ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                          /mod_base= OTHER
/note= "N = 5' biotinylated cytidine'
                                                                                                                                                                                                                                                                                                                                                                                                         Score 19; DB 1; Length 20;
Pred. No. 99;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                 FEATURE:

NAME/KEY:

LOCATION: 1..20

GTHER INFORMATION: /note= "biotinylated Oligo 14ab"
US-08-833-377-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OPERATING SYSTEM: FC-DUS, MS-DUS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
CLASSIFICATION NUMBER: US/08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Stocalla, John R.
REGISTRATION NUMBER: 015389-001610US
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
TEMPRATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Pred. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-838-545-9/c; Sequence 9, Application US/08838545; Patent No. 6046307; Batent INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          361 AGGCCGCAGGAAGAGGAAC 379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20 AGGCCGCAGGAAGAGAAC 2
(415) 576-0200
            TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 14
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                    NAME/KEY: modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                       LOCATION: 1
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                       TOPOLOGY: linear
MOLECULE TYPE: DNA
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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other nucleic acid (PNA),
/desc = "peptide nucleic acid (PNA),
where (deoxy/ribose-phosphate linkages are replaced by
N.(2-aminoethyl)jycine units linked to nucleotide bases via
glycine amino N through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     other nucleic acid (PNA), / desc = "peptide nucleic acid (PNA), where (deoxy(ribose-phosphate linkages are replaced by N-(2-aminoechyl)glycine units linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
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APPLICANT: Shay, Jerry W.
APPLICANT: Wright, Woodring E.
APPLICANT: Wright, Woodring E.
APPLICANT: Wright, Woodring E.
APPLICANT: Orey, David R.
APPLICANT: Corey, David R.
APPLICANT: Or INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Bighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                          4.0%; Score 18; DB 1; Length 18; 100.0%; Pred. No. 1e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 18;
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ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/349,532
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
1e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/838,545
FILING DATE: 09-APR-1997
PILING DATE: 09-APR-1996
ATTONNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,94
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 9, Application US/09349532
Patent No. 6294650
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET WUMBER: 015:3
TELECOMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
                                                                                                                                                                                                                                                                                                        48 AACCCTAACTGAGAAGGG 65
                                                                                                                                                                                                                                                                                                                                                          18 AACCCTAACTGAGAAGGG 1
                                                                                                                                                                                                                      Local Similarity 100.
es 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       single
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3.6%; Score 16.4; DB 1; Length 18; 94.4%; Pred. No. 1.4e+02; ive 0; Mismatches 1; Indels
TITLE OF INVENTION: Human Telomerase Catalytic Subunit WIMBER OF SEQUENCES: 727
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LOCATION: 1..18
OTHER INFORMATION: /note= "antisense hTRT molecule"
                                                                                                                                                                       COUNTRY: CALLLIAGE
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/974,549A
FILING DATE: 19-NOV-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/44,419
FILING DATE: 10-APR-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/851,843
FILING DATE: 25-APR-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/851,843
FILING DATE: 14-AUG-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/851,843
FILING DATE: 14-AUG-1997
PRIOR APPLICATION DATE: US 08/911,312
FILING DATE: 14-AUG-1997
PRIOR APPLICATION DATE: US 08/911,312
FILING DATE: US 08/911,312
FILING DATE: US 08/912,951
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002610US
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0300
INPORMATION FOR SEQ ID NO: 543:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   149 CACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3.6%
Best Local Similarity 94.4%
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: DNA
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          Gaps
                                                                                                                                                                                                                                                                                                                                       APPLICANT: No. 1, Fred APPLICANT: No. 1, Fred APPLICANT: Kealey, James T. APPLICANT: Kealey, James T. APPLICANT: Pruzan, Ronald APPLICANT: Pruzan, Ronald APPLICANT: Weinrich, Scott L. TITLE OF INVENTION: Methods for Detecting the RNA Component of TITLE OF INVENTION: Telomerase NUMBER OF SEQUENCES: 26 CORRESPONDENCE ADDRESSE: TOWNSEND and TOWNSEND and CREW LLP STREET: Two Embarcadero Center, 8th Floor CITY: San Francisco CITY: San Francisco STATE: California
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100.0%; Pred. No. 1.2e+02;
tive 0; Mismatches 0; Indels
        0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: CAILIBETHAN
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/770,565
FILING DATE: 20-DEC-1996
CLASSIFICATION: 435
ATTONREY/AGENT INPORMATION:
NAME: Storella 'John R:
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 31,944
REJERPAK: 415-576-0200
TELEPRAK: 415-576-0300
INPORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
LENGTH: 17 base pairs
        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 156
2.08-97-4549A-543/C
) Sequence 543, Application US/08974549A
) Patent No. 6166178
                                                                                                                                                                                                                                     Sequence 14, Application US/08770565
Patent No. 5846723
GENERAL INFORMATION:
APPLICANT: Kim, Nam Woo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cech, Thomas R.
Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin B.
Andrews, William H.
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                                                           48 AACCCTAACTGAGAAGGG 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                177 TGTCAGCTGCTGGCCCG 193
                                                                                                            18 AACCCTAACTGAGAAGGG 1
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        18; Conservative
                                                                                                                                                                                                                                                                                                                    Kim, Nam Woo
Wu, Fred
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Cech, Th
APPLICANT: Lingner,
APPLICANT: Nakamura
APPLICANT: Chapman,
APPLICANT: Morin, G
APPLICANT: Harley,
APPLICANT: Andley,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           , MOLECULE TYPE: DNA
US-08-770-565-14
                                                                                                                                                                                     RESULT 155
US-08-770-565-14/c
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          Matches
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                Length 18;
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ZIP: 94111-3834

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: EACHLIN PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIN Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/721,456

FILING DATE: 12-NOV-1997

APPLICATION NUMBER: US/08/974,549A

FILING DATE: 19-NOV-1997

APPLICATION NUMBER: US 08/844,419

FILING DATE: 10-NOV-1997

APPLICATION NUMBER: US 08/846,017

FILING DATE: 10-NOV-1997

APPLICATION NUMBER: US 08/846,017

FILING DATE: 06-MAY-1997

APPLICATION NUMBER: US 08/851,843

FILING DATE: 06-MAY-1997

APPLICATION NUMBER: US 08/854,050

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: US 08/912,951

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: US 08/912,951

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: US 08/912,951

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: US 08/915,603

FILING DATE: 11-AUG-1997

APPLICATION NUMBER: US 08/915,603
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Harley, Calvin B.
Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 727
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                             Indels
      3.6%; Score 16.4; DB 1;
94.4%; Pred. No. 1.4e+02;
iive 0; Mismatches 1;
                                                                                                                                                                                                                                                               RESULT 158
US-09-721-456-543/c
; Sequence 543, Application US/09721456
; Patent No. 6617110
. CRNERAL INFORMATION:
. Cech, Thomas R.
                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:

APPLICANT: Cech, Thomas R.

Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                   149 CACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (415) 576-0300
                                                                                                                                                                       18 CACCCTTCATTCTAGAGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 543:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 18 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: San Francisco
STATE: California
Ouery Match
Best Local Similarity 94.4<sup>†</sup>
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: USA
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156 CATTCTAGAGCAAACAAAAA 176
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PCT-US92-10621-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCT-US94-02233-13
                                        임
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Gorman, Cornelia M.,
Groskreutz, Debyra J.
TITLE OF INVENTION: Prohormone Convertase Transformed Cells and
Polypeptide Synthesis
                                                                                                                                                                                                                                                                                                    ö
                                                                                            NOCATION: 1.18
OTHER INFORMATION: /note= "antisense hTRT molecule"
SEQUENCE DESCRIPTION: SEQ ID NO: 543:
US-09-721-456-543
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85.7%; Pred. No. 1.6e+02;
tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                       Length 18
                                                                                                                                                                                                                                                                                                    1; Indels
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Winbatin (Genentech) CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/026,143B FILING DATE: 01-Mar-1993 CLASSFICATION OBTA: APPLICATION NUMBER: 07/803631 FILING DATE: 22-May-1992 APPLICATION NUMBER: 07/803631 FILING DATE: 06-DEC-1992 APPLICATION NUMBER: 07/803631 FILING DATE: 06-DEC-1992 APPLICATION NUMBER: PCT/US92/10621 FILING DATE: 04-DEC-1992 APPLICATION NUMBER: PCT/US92/10621 ATTORNEY/ARTING DATE: 04-DEC-1992 ATTORNEY/ARTING DATE: 04-DEC-1992 ATTORNEY/ARTING DATE: 04-DEC-1992
                                                                                                                                                                                                                                                  Query Match

3.6%; Score 16.4; DB 1;
Best Local Similarity 94.4%; Pred. No. 1.4e+02;
Matches 17; Conservative 0; Mismatches 1;
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REGISTRATION NUMBER: 34,659
REFERENCE/DOCKET NUMBER: P0748P3
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-5530
TELEFA: 650/922-9881
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: Linear SEQUENCE DESCRIPTION: SEQ ID NO: 13: US-08-026-143B-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 13, Application US/08026143B Patent No. 6348327 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: South San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                               149 CACCGTTCATTCTAGAGC 166
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA
PEATURE:
                                                                                                                                                                                                                                                                                                                                                                                            18 CACCCTTCATTCTAGAGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 57
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Best Local Similarity 85.7
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: USA
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Sequence 13, Application PC/TUS9210621
GENERAL INFORMATION:
APPLICANT: Genentech, Inc.
APPLICANT: Gorman, Cornelia M.,
APPLICANT: Marriott, Dave,
APPLICANT: Groskreutz Debrya J.
TITLE OF INVENTION: Prohormone Convertase Transformed Cells and Polypeptide Synthes
NUMBER OF SEQUENCES: 54
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
ADRESSEE: Genentech, Inc.
ATREET: 460 Point San Bruno Blvd
CITY: South San Francisco
STREET: Allfornia
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GENERAL INFORMATION:
APPLICANT: Generatech, Inc.
TITLE OF INVENTION: Prohormone Convertage Transformed Cells and Polypeptide Synthe:
NUMBER OF SEQUENCES: 54
CORRESPONDENCE ADDRESS:
ADDRESSEE: Generatech, Inc.
STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
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85.7%; Pred. No. 1.6e+02;
tive 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: USA
ZIP: 94080
COMPUTER READABLE FORM:
MEDIUM TYPE: 5.25 inch, 360 Kb floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURREATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: patin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/10621
FILING DATE: 19921204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/887265
FILING DATE: 22-MAY-1992
PRIOR APPLICATION DATA:
APPLICATION UMBER: 07/803631
FILING DATE: 06-DEC-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            156 CATTCTAGAGCAAAAAA 176
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CATTCTAGAGCAAAGAGACAA 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Adler, Carolyn R.
REGISTRATION NUMBER: 32,324
REFRENCE/DOCKET NUMBER: 748P
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/952-9614
TELEFAX: 415/952-9681
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 85.7
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 21 bases
TYPE: NUCLEIC ACID
STRANDEDNESS: single
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MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "peptide nucleic acid (PNA),
DESCRIPTION: where (deoxy(ribose-phosphate linkages are replaced by
DESCRIPTION: N-(2-aminoethyl)glycine units linked to nucleotide bases via
DESCRIPTION: glycine amino N through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Shay, Jerry W.
APPLICANT: Wright, Woodring E.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: David R.
APPLICANT: Corey, David R.
APPLICANT: Corey, David R.
APPLICANT: No. 6294650ron, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Bighth Floor CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 3.5%; Score 16; DB 1; Length 16; Best Local Similarity 100.0%; Pred. No. 1.3e+02; Matches 16; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/349,532
             NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPAS: (415) 576-0300
INFORMATION FOR SEQ ID NO: 27:
SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        015389-001610US
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/838,545
FILING DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 27, Application US/09349532
Patent No. 6294650
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TELECOMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
INFORMATION FOR SEQ ID NO: 27:
SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 53 TAACTGAGAAGGGCGT 68
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APPLICANT: Shay, Jerry W.
APPLICANT: Platyszek, Mieczyslaw A.
APPLICANT: Platyszek, Mieczyslaw A.
APPLICANT: Corey, David R.
APPLICANT: No. 6046307ton, James C.
TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS: 60
CORRESPONDENCE ADDRESS: 57
STREET: Townsend and Townsend and Crew LLP
STREET: Townsend and Townsend and Crew LLP
STREET: Townsend and Townsend and Crew LLP
STREET: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3.6%; Score 16.2; DB 1; Length 21; 85.7%; Pred. No. 1.6e+02; tive 0; Mismatches 3; Indels
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APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: 5.25 inch, 360 Kb floppy disk
COMPUTER: IBM PC compatible
COMPUTER: SYSTEM: PC-DOS/MS-DOS
SOFTWARE: patin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/02233
                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
ATOMOSTICATION:
NAME: LOVE, RICHARD B.
REGISTRATION NUMBER: 34,659
REGISTRATION NUMBER: 34,659
RECOMMUNICATION INFORMATION:
TELEPHONE: 415,225-5530
TELEFAX: 415,952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          156 CATTCTAGAGCAAACAAAAA 176
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
CMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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Patent No. 6046307
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAK: 415/952
TELEFAK: 415/952
TELEK: 910/311-7168
; INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 bases
TYPE: nucleic acid
TYPE: --wess: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity 85.7
Matches 18; Conservative
California
: USA
                                                                                                                                                                                                                                                   FILING DATE:
CLASSIFICATION:
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                                                     94080
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PCT-US94-02233-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-838-545-27
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Gaps .. 0

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Gaps
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APPLICANT: Realey, James T.
APPLICANT: Weinrich, Scott L.
TITLE OF INVENTION: Methods for Detecting the RNA Component of TITLE OF INVENTION: Telomerase
NUMBER OF INVENTION: 26
CORRESPONDENCE ADDRESS:
ADDRESSE: TWO Embarcadero Center, 8th Floor
CITY: San Francisco
STATE: California
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                                                                                                                        3.4%; Score 15.2; DB 1; Length 20; 85.0%; Pred. No. 1.8e+02;
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Pred. No. 1.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE PATENTIN Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/770,565
FILING DATE: 20-DEC-1996
CLASSIFICATION: 435
ATTONNEY/AGENT INFORMATION:
NAME: Storella, John R. REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-002300US
TELEPHONIC 415-576-0300
TELEPHONIC: 415-576-0300
                                                                                                                                                                       0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Sequence 12, Application US/08630019A; Patent No. 6015710; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 10, Application US/08770565; Patent No. 5846723; GENERAL INFORMATION:
                                                                                                                                                                                                                           313 CTGTCAGCCGCGGGTCTCTC 332
                                                                                                                                                                                                                                                           CIGICIGCCACGGGTITCIC 20
ORGANISM: Artificial Sequence
                                               ; OTHER INFORMATION: PCR primer US-09-596-938-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    152 CGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 415-576-0300
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.00 Junery Match
Best Local Similarity 100.0
Matches 15, Conservative
                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Kim, Nam Woo APPLICANT: Wu, Fred
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: DNA
                                                                                                                                                  Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 94111-3834
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US-08-770-565-10/c
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                                                                                                                          Query Match
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| Sequence 76, Application US/09345882
| Patent No. 6399373
| GENERAL INFORMATION:
| APPLICANT: Bougueleret, Lydie
| TITLE OF INVENTION: A NUCLEIC ACID ENCODING A RETINOBLASTOMA BINDING PROTEIN (RBP-7)
| TITLE OF INVENTION: AND POLYMORPHIC MARKERS ASSOCIATED WITH SAID NUCLEIC ACID.
| FILE REFERRANCE: GENSET. 031A
| CURRENT FILING DATE: 1999-06-30
| PRIOR APPLICATION NUMBER: US 60/091,315
| PRIOR APPLICATION NUMBER: US 60/091,315
| PRIOR PPLING DATE: 1998-12-10
| PRIOR PILING DATE: 1998-12-10
| PRIOR PLING DATE: 1998-12-10
| NUMBER OF SEQ ID NOS: 140
| SEQ ID NO 76
| LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: primer_bind
| LOCATION: 1..20
| OTHER INFORMATION: upstream amplification primer for SEQ 34, SEQ 55, SEQ 35, SEQ 56
| US-09-345-882-76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Li, Xiao-Jiang
APPLICANT: Li, Xiao-Jiang
APPLICANT: Li, Shi-Una
TITLE OF INVENTION: Huntington Disease Cellular Model:
TITLE OF INVENTION: Stably Transfected FC12 Cells Expressing Mutant Huntingtin
TITLE OF INVENTION: Stably Transfected FC12 Cells Expressing Mutant Huntingtin
TITLE OF INVENTION: Stably Transfected FC12 Cells Expressing Mutant Huntingtin
TITLE OF INVENTION: Stably Transfected FC12 Cells Expressing Mutant Huntingtin
CURRENT APPLICATION NUMBER: US/09/596,938
CURRENT FILING DATE: 2000-06-19
FRIOR APPLICATION NUMBER: 60/140.018
FRIOR PILING DATE: 1999-06-18
NUMBER OF SEQ ID NOS: 15
SOFTWARE: FASTSEQ for Windows Version 3.0
                                            MOLECULE TYPE: other nucleic acid

BESCRIPTION: /desc = "peptide nucleic acid (PNA),

BESCRIPTION: where (deoxy(ribose-phosphate linkages are replaced by

BESCRIPTION: N-(2-aminoethyl)glycine units linked to nucleotide bases via

BESCRIPTION: glycine amino N through a methylenecarbonyl linker"

US-09-349-532-27
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                                                                                                                                                                                                                      Query Match 3.5%; Score 16; DB 1; Length 16; Best Local Similarity 100.0%; Pred. No. 1.3e+02; Matches 16; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1; Indels
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94.1%; Pred. No. 1.7e+02;
tive 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       // Sequence 11, Application US/09596938
// Patent No. 6355481
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  166 CAAACAAAAATGTCAG 182
                                                                                                                                                                                                                                                                                                                              53 TAACTGAGAAGGGCGT 68
                                                                                                                                                                                                                                                                                                                                                                        1 TAACTGAGAAGGGCGT 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA ORGANISM: Artificial Seguence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CAAACAATAAATGTCAG
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Matches 16; Conservative
  single
STRANDEDNESS:
TOPOLOGY: 1i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-596-938-11
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LENGTH: 20
TYPE: DNA
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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Sequence 40, Application US/08630019A

Patent No. 6015710

GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IRM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 01
TELECOMMUNICATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          42 TIGICIAACCCIAAC 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15 rrgrcraacccraac 1
                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: single
              COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear
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where (deoxy)ribose-phosphate linkages are replaced by
M-(2-aminoethyl)glycine units linked to nucleotide bases via
glycine amino nitrogen through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
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US-08-630-019A-18/C
; Sequence 18, Application US/08630019A
; Patent No. 6015710
; GENERAL INFORMATION:
; APPLICANT: Wright, Woodring E.
; APPLICANT: Piatyszek, Mieczyslaw A.
; APPLICANT: Oresy, David
; APPLICANT: No. 6015710ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by
TITLE OF SEQUENCES: 46
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: TOWNSEND and Townsend and Crew LLP
APPLICANT: Shay, Jerry W.
APPLICANT: Wright, Woodring E.
APPLICANT: Wright, Woodring E.
APPLICANT: Corey, David
APPLICANT: Corey, David
APPLICANT: No. 6015710ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by
TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 46
CORRESPONDENCE ADDRESS:
ADDRESSER: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 3.3%; Score 15; DB 1; Length 15; Best Local Similarity 100.0%; Pred. No. 1.4e+02; Matches 15; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                      COUNTRY: USA
ZIP: 94111-3834
COMPUTER FADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/630,019A
FILING DATE: 09-JUN 1996
ATTOREY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001600US
FELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE FHARACTERISTICS:
LENGTH: 15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                other nucleic acid
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DESCRIPTION:
US-08-630-019A-12
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DESCRIPTION:
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MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "peptide nucleic acid (PNA),
DESCRIPTION: where (deoxy)ribose-phosphate linkages are replaced by
DESCRIPTION: where (deoxy)ribose-phosphate linkages are replaced by
DESCRIPTION: N-(2-aminoethyl)iglycine units linked to nucleotide bases via
DESCRIPTION: glycine amino nitrogen through a methylenecarbonyl linker"
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APPLICANT: Wright, Woodring E.
APPLICANT: Wright, Woodring E.
APPLICANT: Corey, David
APPLICANT: Corey, David
APPLICANT: No. 6015710ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by
TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 46
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
SOFTWARES PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/630,019A
FILING DATE: 09-JUN 1996
CLASSIFICATION: 536
ATTONREY/AGENT INFORMATION:
NAME: Storella, John R:
REGISTRATION NUMBER: 32,944
REFRENCE/POCKET NUMBER: 32,944
REPRENCE/POCKET NUMBER: 32,944
REPRENCE/POCKET NUMBER: 32,944
REPRENCE/POCKET NUMBER: 32,944
REPRENCE/POCKET NUMBER: 31,944
RELEPHONE: (415) 576-0300
INPORTATION CALLED STORED NO: 18:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
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3.3%; Score 15; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 1.4e+02;
Matches 15; Conservative 0; Mismatches 0;
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Best Local Similarity 100 Matches 15; Conservative
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/desc = "peptide nucleic acid (PNA),
where (deoxy/tibose-phosphate linkages are replaced by
N.(2-aminoethy/)ajvcine units linked to nucleotide bases via
glycine amino N through a methylenecarbonyl linker"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 2, Application US/08838545
; Sequence 2, Application US/08838545
; Patent No. 6046307
; GENERAL INFORMATION:
    APPLICANT: Wright, Woodring E.
    APPLICANT: Piatyszek, Mieczyslaw A.
    APPLICANT: Orey, David R.
    APPLICANT: No. 6046307ton, James C.
    TITLE OF INVENTION: Modulation of Mammalian Telomerase by ITILE OF INVENTION: Peptide Nucleic Acids
    NUMBER OF SEQUENCES: 60
    CORRESPONDENCE ADDRESS: 6
    CORRESPONDENCE ADDRESS: 7
    ADDRESSEE: Townsend and Townsend and Crew LLP
    STREET: Two Embarcadero Center, Eighth Floor
    CITY: San Francisco
    STARET: California
                                                                                LENGTH: 15 base pairs

TYPE: nucleic acid

TYPE: nucleic acid

TYPOLOGY: linear

MOLECULE TYPE: other nucleic acid

DESCRIPTION: /desc = "phosphorothioate (PS) nucleic acid"
                                                                                                                                                                                                                                                                                                3.3%; Score 15; DB 1; Length 15; 100.0%; Pred. No. 1.4e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OPERATING SYSTEM: PC-LUCS/MS-LUCS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 015389-001610US
TELECOMMUNICATION INFORMATION:
SEQUENCE CHARACTERICATICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Palace
                                                                                                                                                                                                                                                                                                                                                                                                            46 CTAACCCTAACTGAG 60
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      15 CTAACCCTAACTGAG 1
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TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: other nucl
DESCRIPTION: /desc = "p
                                                                                                                                                                                                                                                                                                                                                       15; Conservative
                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 15; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
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US-08-838-545-2/c
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DB 1; Length 15;

3.3%; Score 15;

Query Match

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/desc = "peptide nucleic acid (PNB), where (deoxy/tibose-phosphate linkages are replaced by NR.(2-aminoethyl)glycine units linked to nucleotide bases vis glycine amino N through a methylenecarbonyl linker"
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                                           Gaps
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                                                                                                                                                                                                                                                                                                  APPLICANT: Shay, Jerry W.
APPLICANT: Wright, Woodring E.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Diatyszek, Mieczyslaw A.
APPLICANT: Corey, David R.
APPLICANT: No. 6046307ton, James C.
APPLICANT: No. 6046307ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by
TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3.3%; Score 15; DB 1; Length 15; 100.0%; Pred. No. 1.4e+02; tive 0; Mismatches 0; Indele
                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: FLORDY disk
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
CORRENTING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/818,545
FILING DATE: 09-APR-1997
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: SECOREL NUMBER: 015389-001610US
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARATERISTICS:
                  Pred. No. 1.4e+02;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   other nucleic acid
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100.0%; Pre-
                                                                                                                                                                                                                                      ; Sequence 5, Application US/08838545; Patent No. 6046307
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                                                                                                                           15 craacccraacreae 1
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Best Local Similarity 100.0
Matches 15, Conservative
                                                                                  46 CTAACCCTAACTGAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: San Francisco
STATE: California
COUNTRY: USA
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DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
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STRANDEDNESS:
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DESCRIPTION:
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Sequence 2, Application US/09349532

Patent No. 6294650

GENERAL INPORMATION:
APPLICANT: Shay, Jarry W.
APPLICANT: Wright, Woodring E.
APPLICANT: Wright, Woodring E.
APPLICANT: Diatyszek, Mieczyslaw A.
APPLICANT: Orey, David R.
APPLICANT: No. 6294650ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS: 7
CORRESPONDENCE ADDRES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3.3%; Score 15; DB 1; Length 15; 100.0%; Pred. No. 1.4e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                               CITY: San Francisco Center, Eighth Floor STATE: California COUNTRY: USA ZIP: 94111-3834
COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBM PC Compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/349,532
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "phosphorothioate (PS)

JESCRIPTION: nucleic acid"
US-08-838-545-45
                                                                                          COUNTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk

"MEDUWER: IBM PC compatible
""GTEM: PC-DOS/MS-DOS
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Best Local Similarity 100.0
Matches 15; Conservative
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                           San Francisco
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PRIOR APPLICATION DATA:
                                                          California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
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US-09-349-532-2/c
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                   APPLICANT: Shay, Jerry W.
APPLICANT: Wright, Woodring E.
APPLICANT: Platyszek, Mieczyslaw A.
APPLICANT: Corey, David Meczyslaw A.
APPLICANT: No. 6046307ton, James C.
APPLICANT: No. 6046307ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 45, Application US/08838545

Patent No. 6046307

GENERAL INFORMATION:
APPLICANT: Wright, Woodring E.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Corey, David R.
APPLICANT: No. 6046307ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60

CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 15;
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                                                                                                                                                                                                                                                                                                                                             CURKES-CONDENCE AUDKESS:
ADDREESSEE: Two Embarcadero Center, Eighth Floor
STREET: Two Embarcadero Center, Eighth Floor
STRATE: California
COUNTRY: USA
ZIP: 9411-3834
COMPUTER: TEM PC Compatible
COMPUTER: TEM PC Compatible
COPERATING SYSTEM: PC-DOS/MS-DOS
CURENT APPLICATION DATA:
APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
CLASSIFICATION DATA:
APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
CLASSIFICATION DATA:
APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
FILING DATE: 09-APR-1997
CLESSIFICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
FILING DATE: 09-APR-1997
CLESSIFICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1996
FILING DATE: 09-APR-1997
FILING DATE: 09-APR-1996
FILING DATE: 09-APR-1997
FILING DATE: 09-APR-1996
FILING DATE: 09-APR-1996
FILING DATE: 09-APR-1997
FILING DATE: 09-APR-199
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Pred. No. 1.4e+02;
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Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 3.3%; Score 15; DB Best Local Similarity 100.0%; Pred. No. 1.4 Matches 15; Conservative 0; Mismatches
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: lin
MOLECULE TYPE:
DESCRIPTION:
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DESCRIPTION:
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US-08-838-545-45/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-838-545-28
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Gaps

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other nucleic acid / desc = "poptide nucleic acid (PNA), where (deoxy(ribose-phosphate linkages are replaced by N-(2-aminoethyl)glycine units linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Shay, Jerry W. APPLICANT: Wight, Woodring E. APPLICANT: Wright, Woodring E. APPLICANT: Platyszek, Misczyslaw A. APPLICANT: Corey, David R. APPLICANT: Corey, David R. TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids NUMBER OF SEQUENCES: 60 CORRESPONDENCE ADDRESS: ADDRESSER: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                       Length 15;
                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/349,532
FILING DATE:
CLASSIFICATION
PRIOR APPLICATION
APPLICATION DATA:
PRIOR APPLICATION
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REFERENCE/POCKET NUMBER: 32,944
REFERENCE/POCKET NUMBER: 32,944
REFERENCE/POCKET NUMBER: 015389-001610US
                                                                                                                                                                                                                                                                                                                                       Query Match
3.3%; Score 15; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 1.4e+02;
Matches 15; Conservative 0; Mismatches 0;
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Patent No. 6294650
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                   42 TIGICTAACCCTAAC 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
         TELEFAX: (415) 576-0300 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (415) 576-030
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                           LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM:
                                                                                                                                                      linear
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                                                                                                                                                 TOPOLOGY: lin
MOLECULE TYPE:
DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
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DESCRIPTION:
DESCRIPTION:
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                                                                                                                                                                                                                                                                                                                                                                                                          DESCRIPTION: /desc = "peptide nucleic acid (PNA),
DESCRIPTION: /desc = "peptide nucleic acid (PNA),
DESCRIPTION: where (deoxy(ribose-phosphate linkages are replaced by
DESCRIPTION: where ylloglycine units linked to nucleotide bases via
DESCRIPTION: glycine amino N through a methylenecarbonyl linker"
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APPLICATION NUMBER: US/09/349,532
APPLICATION NUMBER: US 08/838,545
FILING DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELEPAK: (415) 576-0200
TELEPAK: (415) 576-0200
TELEFAK: (415) 576-0300
TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "poptide nucleic acid (
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CLASSIFICATION DATA:
PRIOR PAPLICATION DATA:
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELECOMMUNICATION INPORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      46 CTAACCCTAACTGAG 60
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Best Local Similarity 100.
Matches 15; Conservative
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JS-09-673-298-4/C
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N-(2-aminoethyl)glycine units linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
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                                                                                                                                                                                                                                                                                                                            Sequence 45, Application US/09349532;
Sequence 45, Application US/09349532;
Batent No. 6294650
GENERAL INFORMATION:
APPLICANT: Wright, Woodring E.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Corey, David R.
APPLICANT: No. 6294650ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STARE: California
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                                                                                          Query Match 3.3%; Score 15; DB 1; Length 15; Best Local Similarity 100.0%; Pred. No. 1.4e+02; Matches 15; Conservative 0; Mismatches 0; Indels
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3.3%; Score 15; DB 1; Length 15;
Best Local Similarity 100.0%; Pred. No. 1.4e+02;
Matches 15; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION NATR:
APPLICATION NUMBER: US/09/349,532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
CLASSIFICATION
PRIOR APPLICATION
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELEPHONE: (415) 576-020
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 45:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                other nucleic acid
/desc = "phosphorothioate (PS)
nucleic acid"
                                                                                                                                                                                    49 ACCCTAACTGAGAAG 63
                                                                                                                                                                                                                          1 Acceraacreagaag 15
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    DESCRIPTION:
    DESCRIPTION:
    US-09-349-532-28
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DESCRIPTION:
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other nucleic acid
/desc = "peptide nucleic acid (PNA),
where (deoxy)ribose-phosphate linkages are replaced by
N-(2-aminoethyl)alycine units linked to nucleotide bases via
glycine amino nitrogen through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 3.1%; Score 14; DB 1; Length 14; Best Local Similarity 100.0%; Pred. No. 1.6e+02; Matches 14; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       us-108-158-352-2
; Sequence 2, Application US/08158352
; Patent No. 5700922
; GENERAL INFORMATION:
; APPLICANT: Philip Dan Cook
; TITLE OF INVENTION: Macromolecules
; TITLE OF INVENTION: Macromolecules
; TORRESPONDENCES: 4
; CORRESPONDENCES: 4
; ADDRESSEE: No. 5700922xi8
; ADBRESSEE: No. 5700922xi8
; STREET: One Liberty Place - 46th Floor
CITY: Philadelphia
CONTREY: U.S. b
ZIP.
                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATE:
APPLICATION NUMBER: US/08/630,019A
FILING DATE: 09-JUN-1996
CLASSIFICATION: 536
ATTORNEY AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 015389-001600US
TELEPAN: (415) 576-020
TELEPAN: (415) 576-020
INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
LENGTH: 14 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
TYPE: TOWNEDDNESS: single
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
                                               Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy disk, 1.44 Mb storage COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Wordberfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/158,352
FILING DATE: herewith
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US92/11339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 52 CTAACTGAGAAGGG 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: Two bulled CITY: San Francisco CTATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
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DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-630-019A-25
                                                                                           STATE: Ca
COUNTRY:
ZIP: 9411
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: Description of Artificial Sequence: antisense OTHER INFORMATION: oligonucleotide corresponding to a specific region OTHER INFORMATION: of the mouse Ii gene. US-09-205-295-24
                                                                                                                                                                                                                                                                                                                                                                                    ö
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                                                                                                                                                                                                                                                                                                                                   Query Match 3.3%; Score 14.8; DB 1; Length 19; Best Local Similarity 88.9%; Pred. No. 1.8e+02; Matches 16; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Shay, Jerry W.
APPLICANT: Wright, Woodring E.
APPLICANT: Wright, Woodring E.
APPLICANT: Platyszek, Micczyslaw A.
APPLICANT: Corey, David
APPLICANT: No. 6015710ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by
TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Xu, Minzhen
APPLICANT: Xu, Gang
APPLICANT: Yu, Gang
APPLICANT: Humphreys, Robert
ITILE CANT: Humphreys, Robert
ITILE OF INVENTION: CANCER CELL VACCINE
CURRENT APPLICATION NUMBER: 03/09/205,995
CURRENT FILING DATE: 1998-03-09
PRIOR APPLICATION NUMBER: 09/036,746
PRIOR APPLICATION NUMBER: 09/61,627
PRIOR APPLICATION NUMBER: 09/661,627
RIOR APPLICATION NUMBER: 1996-06-11
NUMBER OF SEQ ID NOS: 79
SEQ ID NO S: 79
SEQ ID NO S: 79
LENGTH: 18
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US-08-630-019A-25/c
; Sequence 25, Application US/08630019A
; Patent No. 6015710
                                                                                                               TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "PRIMER"
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 24, Application US/09205995; Patent No. 6368855; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                 348 GTTCAGGCCTTTCAGGCC 365
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 GATCAGGCCTTTTAGGCC 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 GGTGGCCTGCCCAGCC 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA ORGANISM: Artificial Sequence
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
                                                                                                                                                                                                                                                                Homo sapiens
                                                                     TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                    ORIGINAL SOURCE:
ORGANISM: Hom
                                                                                                                                                                                                                ANTI-SENSE: NO
                                                                                                                                                                                       HYPOTHETICAL:
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Zwick, Michael G.
Edington, Brent E.
McSwiggen, James A.
McHo, Patricia Ann Owens
Guo, Lining
Skokut, Thomas A.
Young, Scott A.
Folkerts, Otto
      TITLE OF INVENTION: MODULATION OF TITLE OF INVENTION: IN PLANTS
NUMBER OF SEQUENCES: 1263
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 571, Application US/08679645
Patent No. 6350934
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             106 CGCTGACTTTCAGCGGG 122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity 64.7
Matches 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                  CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
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                                                                                                                                                                                                                                                                                                                                                                                    Query Match 3.1%; Score 13.8; DB 1; Length 17; Best Local Similarity 88.2%; Pred. No. 1.9e+02; Matches 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 2, Application US/09108911

Patent No. 6277603

GENERAL INFORMATION:

APPLICANT: Cook, Phillip Dan

TILE OF INVENTION: PNA-DNA-DNA Chimeric Macromolecules

FILE OF INVENTION: B1S3102

CURRENT APPLICATION NUMBER: US/09/108,911

CURRENT APPLICATION NUMBER: US/09/108,911

CURRENT APPLICATION NUMBER: 08/877,317

PRIOR PILING DATE: 1997-06-17

NUMBER: OF SEQ ID NOS: 4

SOFTWARE: Patentin version 3.0

LENTH: 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Zwick, Michael G.
APPLICANT: Edington, Brent E.
APPLICANT: McSwiggton, James A.
APPLICANT: McSwiggton, James A.
APPLICANT: Merlo, Patricia Ann Owens
APPLICANT: Skokut, Thomas A.
APPLICANT: Skokut, Thomas A.
APPLICANT: Polkerts, Otto
APPLICANT: Polkerts, Otto
APPLICANT: Merlo, Donald J.
TITLE OF INVENTION: COMPOSITION AND METHODS FOR
                                                                                               ISIS-1236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; NAME/KEY: misc_feature
; OTHER INFORMATION: Antisense Sequence
US-09-108-911-2
FILING DATE: December 23, 1992
ATTORNEY/AGENT INFORMATION:
NAME: JOHN W. CAIDWALL
REGISTRATION NUMBER: 28,937
REFERENCE/DOCKET NUMBER: 151S-12:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-568-3100
TELEPAX: 215-568-3439
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Sequence 826, Application US/08679645; Patent No. 6350934; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          102 TTCTCGCTGACTTTCAG 118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TTCTCGCTGACTTTCAG 118
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
ANTI-SENSE: yes
                                                                                                                                                                                                                               LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 184
US-08-679-645-826
                                                                                                                                                                                                                                                                                                                                          US-08-158-352-2
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Edington, Brent G.
APPLICANT: Edington, Brent E.
APPLICANT: McSwiggen, James A.
APPLICANT: McNo, Juning
APPLICANT: Guo, Lining
APPLICANT: Skokut, Thomas A.
APPLICANT: Skokut, Thomas A.
APPLICANT: Skokut, Thomas A.
APPLICANT: Foung, Scott A.
APPLICANT: Poung, Scott A.
APPLICANT: Mclo, Donald J.
TITLE OF INVENTION: MODULATION OF GENE EXPRESSION
TITLE OF INVENTION: IN PLANTS
NUMBER OF SEQUENCES: 1263
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: Suite 4700
MODULATION OF GENE EXPRESSION
IN PLANTS
1263
                                                                                                                                                                                                                                                                                                    LIF: 900/1-2006
COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: WORD PEFFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/679,645
FILING DATE: July 12, 1996
CLASSIFICATION NUMBER: US/08/679,645
FILING DATE: July 12, 1996
RICA APPLICATION NUMBER: 00/001,135
APPLICATION NUMBER: 00/001,135
APPLICATION NUMBER: 00/3100,726
FILING DATE: September 2, 1994
ATTORNEY AGENT INFORMATION:
NAME: WAYDURY RICHARD J.
REGISTRATION NUMBER: 32,327
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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US-10-232-634-11
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; Sequence 11, Application US/09586376
; Patent No. 6492115
; GENERAL INFORMATION:
; APPLICANT: Guida, Marco
; APPLICANT: Hall, Jeff
; TITLE OF INVENTION: GENETIC TYPING OF THE HUMAN CYTOCHROME P450 2A6 GENE
; TITLE OF INVENTION: AND RELATED MATERIALS AND METHODS
; FILE REFRENCE: 4389-20
; CURRENT PAPLICATION WUMBER: US/09/586,376
; CURRENT FILING DATE: 2000-06-02
; NUMBER OF SEQ ID NOS: 29
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 11
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Pred. No. 2e+02;
0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7. Match 3.1%; Score 13.8; D. Local Similarity 82.4%; Pred. No. 2e+0 1es 14; Conservative 1; Mismatches
                               STATE: CALLLOLLE
COUNTRY: U.S.A..
ZIP: 90071-2066

ENDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: BECOMPATER: IBM COMPATIBLE OCHORATION DATA: APPLICATION NUMBER: US/08/679,645 FILING DATE: July 12, 1996
CLASSIFICATION NUMBER: 00/001,135 FILING DATE: September 2, 1994
APPLICATION NUMBER: 00/001,135 FILING DATE: September 2, 1994
ATTORNEY/AGENT INFORMATION: MAME: Warburg, Richard J. REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 219/247
TELECOMMUNICATION INFORMATION: TELEFONDER: (3.13) 955-0440
TELEFONE: TELEFONDE: (3.13) 955-0440
TELEFONE: SEGURACE CHARACTERIES: SEGURACE THEORY SEGURACE CHARACTERIES: SEGURACE THEORY SEGURACE CHARACTERIES: SEGURACE THEORY SEGURACE SEGURACE SEGURACE SEGURACE THEORY SEGURACE SEGURACE
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Best Local Similarity 93.3%;
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Los Angeles
California
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Matches
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Sequence 4559, Application US/09371772B

Fatent No. 6566127

GENERAL INFORMATION:
APPLICANT: Ravco, Pan
APPLICANT: Ravco, Pan
APPLICANT: Bavco, Pan
APPLICANT: Becobedo, Jaime
APPLICANT: Escobedo, Jaime
APPLICANT: Becobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Method and Reagent Endothelial Growth Factor Receptor
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHBDO, 876-J (237/198)
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: PatentIn Version 3.0
SEQ ID NO 4559
LENGTH: 17
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Patent No. 6492115
GENERAL INFORMATION
APPLICANT: Hall, Jeff
TITLE OF INVENTION: GENETIC TYPING OF THE HUMAN CYTOCHROME.P450 2A6 GENE
TITLE OF INVENTION: GENETIC TYPING OF THE HUMAN CYTOCHROME.P450 2A6 GENE
TITLE OF INVENTION: AND RELATED MATERIALS AND METHODS
FILE REFERENCE: 4389-20
CURRENT FILING DATE: 2000-06-02
NUMBER OF SEQ ID NOS: 29
SOFTWARE: Patentin Ver. 2.1
LENGTH: 17
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Patent No. 6797477

GENERAL INFORMATION:
APPLICANT: Guida Marco
APPLICANT: Hall, Jeff
TITLE OF INVENTION: GENETIC TYPING OF THE HUMAN CYTOCHROME P450 2A6 GENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gape
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      3.0%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 93.3
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                        ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-586-376-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; ORGANISM: Homo sapiens
US-09-371-772B-4559
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
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Best Local Similarity
Matches 13; Conserv
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STATE: Ca
COUNTRY:
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APPLICANT: Guida, Marco
APPLICANT: Guida, Marco
APPLICANT: Guida, Marco
APPLICANT: Guida, Marco
APPLICANT: Hall, Jeff
TITLE OF INVENTION: GENETIC TYPING OF THE HUMAN CYTOCHROME P450 2A6 GENE
TITLE OF INVENTION: AND RELATED MATERIALS AND METHODS
TITLE OF INVENTION: AND RELATED MATERIALS AND METHODS
FILE REFERENCE: 4389-20
CURRENT APPLICATION NUMBER: US/10/232,634
CURRENT FILING DATE: 2002-06-20
PRIOR APPLICATION NUMBER: US/09/586,376
NUMBER OF SEQ ID NOS: 29
SOFTWARE: PATENTIN VEF. 2.1
SEQ ID NOS: 29
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APPLICANT: Wight, Woodring E.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Corey, David
APPLICANT: No. 6015710ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by
TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 46
CORRESPONDENCE ADDRESS: 46
CORRESPONDENCE ADDRESS: ADDRESSE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                     Query Match
3.0%; Score 13.4; DB 1; Length 17;
Best Local Similarity 93.3%; Pred. No. 2e+02;
Matches 14; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3.0%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 2e+02; Live 0; Mismatches 1; Indels
TITLE OF INVENTION: AND RELATED MATERIALS AND METHODS FILE REFERENCE: 4389-20
CURRENT APPLICATION NUMBER: US/10/232,634
CURRENT FILING DATE: 2002-08-30
PRIOR APPLICATION NUMBER: US/09/586,376
PRIOR FILING DATE: 2000-06-02
NUMBER OF SEQ ID NOS: 29
SOFTWARE: PATENTIN Ver. 2.1
LENGTH: 17
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; Sequence 11, Application US/08630019A
Patent No. 6015710
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 190
US-10-232-634-12
Sequence 12, Application US/10232634
Patent No. 6797477
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Best Local Similarity 93.3
Matches 14; Conservative
                                                                                                                                                                                                                                                           ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                      US-10-232-634-11
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US-10-232-634-12
                                                                                                                                                                                                                                            TYPE: DNA
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Length 13;
        DB 1; Lens.
7. 1.7e+02;
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Conservative 0; Mismatches
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                                          Score 13;
Pred. No.
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; Sequence 1, Application US/08838545
; Patent No. 6046307
; GENERAL INFORMATION:
APPLICANT: Shay, Jerry W.
; APPLICANT: Wright, Woodring E.
; APPLICANT: Piatryszek, Mieczyslaw A.
APPLICANT: COREY, David R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-630-019A-41/c
; Sequence 41, Application US/08630019A
Patent No. 6015710
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
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Best Local Similarity 100.
Matches 13; Conservative
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Query Match
Best Local Similarity
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/desc = "peptide nucleic acid (PNA),

where (descy)ribose-phosphate linkages are replaced by

N.(2-aminoethyl)sylvine units linked to nucleotide bases via

glycine amino nitrogen through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                                                                                                                       /desc = "peptide nucleic acid (PNA), where (deoxy)ribose-phosphate linkages are replaced by N-(2-aminoethyl)glycine units linked to nucleotide bases via glycine amino nitrogen through a methylenecarbonyl linker"
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APPLICANT: Wight, Woodring E.
APPLICANT: Wistyle, Woodring E.
APPLICANT: Wistyle, Wieczyslaw A.
APPLICANT: Corey, David
APPLICANT: Corey, David
APPLICANT: No. 6015710ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by
TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 46
CORRESPONDENCES. 46
CORRESPONDENCE ADDRESSE
ADDRESSE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
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MEDIUM TYPE: Floppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/630,019A
FILING DATE: 09-JUN-1996
CLASSIFICATION: 536
ATTORNEY AGENT INFORMATION:
NAME: Storella, John R.
REFERENCE/DOCKET NUMBER: 015389-001600US
TELEPHONE: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 2.9%; Score 13; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 1.7e+02;
Matches 13; Conservative 0; Mismatches 0;
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; Sequence 17, Application US/08630019A
Patent No. 6015710
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                     other nucleic acid
                                      TELEPHONE: (415) 576-0200
TELEPAX: (415) 576-0300
INPORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GICTAACCCTAAC 56
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| MOLECULE TYPE: of DESCRIPTION: |
| DESCRIPTION: of DESCRIPTIO
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MOLECULE TYPE:
DESCRIPTION:
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DESCRIPTION:
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                                                                                                                                                                                                                                                                                                                                  APPLICANT: Shay, Jerry W.
APPLICANT: Widght, Woodring E.
APPLICANT: Play, Mecayslaw A.
APPLICANT: Playsek, Mieczyslaw A.
TITLE OF INVENTION: Modilation of Mammalian Telomerase by
TITLE OF INVENTION: Peptide Nucleic Acids
TITLE OF INVENTION: Peptide Nucleic Acids
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "phosphorothioate (PS) nucleic acid"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2.9%; Score 13; DB 1; Length 13;
100.0%; Pred. No. 1.7e+02;
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CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 09-JUN-1996
CLASSIFICATION:
SAGNATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: SLOCATION: NAME: 32,944
REFERENCE/DOCKET NUMBER: 015389-001600US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 41:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
```

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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATE: US/08/838,545
FILING DATE: 09-APR-1997
RIOR APPLICATION DARR: US/08/838,545
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: SLORella, John R.
REPERENCE/DOCKET NUMBER: 015389-001610US
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 base pairs
TYPE: NUCLEIC acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      other nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   42 TTGTCTAACCCTA 54
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               linear
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MOLECULE TYPE:
DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-838-545-4
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APPLICANT: Shay, Jerry W.
APPLICANT: Shay, Jerry W.
APPLICANT: Platyscak, Mieczyslaw A.
APPLICANT: Platyscak, Mieczyslaw A.
APPLICANT: Corey, David R.
APPLICANT: Corey, David R.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
APPLICANT: No. 6046307ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by
TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/838,545
FLING DATE: 09-APR-1997
CLEASIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY AGENT INFORMATION:
NAME: SLOCKET NUMBER: 32,944
REPERENCE/DOCKET NUMBER: 015389-001610US
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 base pairs
TYPE: MOLDING SIGILS
TYPE: MOLDING SIGILS
TYPE: MOLDING SIGILS
TYPE: MOLDING SIGILS
TYPE: MOLDINGS SIGILS
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2.9%; Score 13; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 1.7e+02;
Matches 13; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       other nucleic acid
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COMPUTER READABLE FORM:
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MOLECULE TYPE:
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DESCRIPTION:
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US-08-838-545-4/c
                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
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/desc = "peptide nucleic acid (PNA),
where (deoxy/tibose-phosphate linkages are replaced by
N-(2-aminoethyl)glycine units linked to nucleotide bases via
glycine amino N through a methylenecarbonyl linker"
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                                                                                                                                                                                                                                                           Gaps
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APPLICANT: Shay, Jerry W.
APPLICANT: Shay, Jerry W.
APPLICANT: Biatyszek, Mieczyelaw A.
APPLICANT: Diatyszek, Mieczyelaw A.
APPLICANT: Corey, David R.
APPLICANT: Corey, David R.
APPLICANT: No. 6046307ton, James C.
ITILE OF INVENTION: Modulation of Mammalian Telomerase by
ITILE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Bighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                      Query Match 2.9%; Score 13; DB 1; Length 13; Best Local Similarity 100.0%; Pred. No. 1.7e+02; Matches 13; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: BATEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 536
PRIOR APPLICATION UNBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
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other nucleic acid

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13 rrrrrrrrrraki
                           ) DESCRIPTION:
) DESCRIPTION:
US-08-838-545-46
       MOLECULE TYPE:
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                                                                                                                                                                        Matches
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where (deoxy/tibose_phosphate linkages are replaced by
N-(2-aminoethyl) alycine units linked to nucleotide bases via
glycine amino N through a methylenecarbonyl linker"
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APPLICANT: Wright, Woodring E.
APPLICANT: Wright, Woodring E.
APPLICANT: Platyszek, Mieczyslaw A.
APPLICANT: Corey, David M.
APPLICANT: Corey, David M.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Modulation of Mammalian Telomerase by NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.9%; Score 13; DB 1; Length 13; 100.0%; Pred. No. 1.7e+02; tive 0; Mismatches 0; Indels
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COUNTER READABLE FORM
ZIP: 94111-3834

ZIP: 94111-3834

COMPUTER READABLE FORM
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 015389-001610US
TELEPAN: (415) 576-0200
TELEPAN: (415) 576-0200
TELEPAN: (415) 576-0300
INFORMATION FOR SEQ ID NO: 46:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELECOMMUNICATION INFORMATION:
TELEPAX: (415) 576-0300
TELEPAX: (415) 576-0300
TELEPAX: GISS 576-0300
TYPE: Nucleic acid STRANDEDNESS: single STRANDEDNESS: single TYPE: nucleic acid STRANDEDNESS: single TYPE: Other nucleic acid DESCRIPTION: /desc = "peptide nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 198
US-08-838-545-46/c
; Sequence 46, Application US/08838545
; Patent No. 6046307;
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         44 GTCTAACCCTAAC 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13; Conservative
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Matches 13; Conserva
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DESCRIPTION:
DESCRIPTION:
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/desc = "peptide nucleic acid (PNB), where (deoxy/ribose-phosphate linkages are replaced by where (deoxy/ribose-phosphate linked to nucleotide bases via 91ycine amino N through a methylenecarbonyl linker"
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                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Shay, Jerry W.
APPLICANT: Wright, Woodring E.
APPLICANT: Wright, Woodring E.
APPLICANT: Piaryszek, Mieczyslaw A.
APPLICANT: Crewy, David R.
APPLICANT: No. 6046307ton, James C.
APPLICANT: No. 6046307ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by
TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
                                                                                               DB 1; Length 13;
1.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2.9%; Score 13; DB 1; Length 13; 100.0%; Pred. No. 1.7e+02; Live 0; Mismatches 0; Indels
                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN RAIS
CORRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/836,545
FILING DATE: 09-APR-1997
CLASSIFICATION NUMBER: US/08/630,019
FILING DATE: 09-APR-1996
PRING APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: SCOTCALIA, JOHN R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0300
INFORMATION INFORMATION:
TELEPHONE: (415) 576-0300
INFORMATION INFORMATION:
TELEPHONE: US SEQUENCE CHARACTERISTICS:
LENGTH: 13 base pairs
TYPE: uncleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
/desc = "phosphorothioate (PS) nucleic acid"
                                                                                                                                               0; Mismatches
                                                                                             2.9%; Score 13;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Townsend and Townsend and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           other nucleic acid
                                                                                                                                                                                                                                                                                                                                                                 Sequence 51, Application US/08838545
Patent No. 6046307
GENERAL INFORMATION:
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Best Local Similarity
                                                                                                                         Best Local Similarity
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DESCRIPTION:
DESCRIPTION:
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TOPOLOGY: linear
MOLBCULE TYPE: other nucleic acid (PNA),
DESCRIPTION: /desc = "peptide nucleic acid (PNA),
DESCRIPTION: where (decoy(ribose-phosphate linkages are replaced by
DESCRIPTION: where (decoy(ribose-phosphate linkages are replaced by
DESCRIPTION: N-(2-aminoethyl)glycine units linked to nucleotide bases via
DESCRIPTION: glycine amino N through a methylenecarbonyl linker"
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NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Bighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 13;
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                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/838,545

FILING DATE: 09-APR-1997

CLASSIFICATION NUMBER: US 08/630,019

FILING DATE: 09-APR-1996

ATTORNEY/AGENT INFORMATION:

NAME: Storella, John R.

REGISTRATION NUMBER: 32,944

REGISTRATION NUMBER: 32,944

REGISTRATION NUMBER: 35.6-0200

TELEPHONE: (415) 576-0300

INFORMATION FOR SEQ ID NO: 55:

SEQUENCE CHARACTERISTICS:

LEMGTH: 13 base pairs

LEMGTH: 13 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2.9%; Score 13; DB 1; Le
100.0%; Pred. No. 1.7e+02;
tive 0; Mismatches 0;
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San Francisco
California
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          53 TAACTGAGAAGGG 65
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
nes 13; Conserv
                                                                                                                                                                                                                        STATE: Ca
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Matches
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/desc = "peptide nucleic acid (PNA),
where (deoxy/tribose-phosphate linkages are replaced by
N.(2-aminoethy/) glycine units linked to nucleotide bases via
glycine amino N through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
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0
                                                                                                                     Sequence 52, Application US/08838545
Patent No. 6046307
GENERAL INPORMATION:
APPLICANT: Shay, Jerry W.
APPLICANT: Carey, Wieczyslaw A.
APPLICANT: Corey, David R.
APPLICANT: Corey, David R.
APPLICANT: No. 6046307ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.9%; Score 13; DB 1; Length 13; 100.0%; Pred. No. 1.7e+02; tive 0; Mismatches 0; Indels
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CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGBNT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELECOMMUNICATION INFORMATION:
TELEBEHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 52:
SEQUENCE CHARACTERICAS:
CHARA
                                                                                                                                                                                                                                                                                                                                                                                                       STREET: Two Embarcadero Center, Eighth Floor STATE: California
UNITRY: USA
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US-08-838-545-55/C
; Sequence 55, Application US/08838545
; Patent No. 6046307
; GENERAL INFORMATION:
APPLICANT: Shay, Jerry W.
APPLICANT: Wright, Woodring E.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Corey, David R.
APPLICANT: No. 6046307ton, James C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPPERATING SYSTEM: PC-DOS/MS-DOS
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Best Local Similarity luv...
Thea 13; Conservative
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TYPE: nucleic acid
STRANDEDNESS: single
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DESCRIPTION:
DESCRIPTION:
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DESCRIPTION:
                                                                         RESULT 200
US-08-838-545-52/c
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Gaps

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/desc = "peptide nucleic acid (PNB), where (deoxy/trbose-phosphate linkages are replaced by NF.(2-aminoethyl)glycine units linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.9%; Score 13; DB 1; Le
100.0%; Pred. No. 1.7e+02;
tive 0; Mismatches 0;
                             NAME: Scorella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELEFONMUNICATION INPORMATION:
TELEFONMUNICATION 10000
TELEFAX: (415) 576-0300
INPORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 base pairs
TYPE: nucleic acid
STRANDENNESS: single
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REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/838,545
FILING DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                  other nucleic acid
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Patent No. 6294650
GENERAL INFORMATION:
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TELEPHONE: (415) 576-0200
TELEPAK: (415) 576-0300
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
          ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity 100.
nes 13; Conservative
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: USA
                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
MOLECULE TYPE: other
                                                                                                                                                                                                                                                                                                                                                                       DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
DESCRIPTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-349-532-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      other nucleic acid
/desc = "peptide nucleic acid (PNA),
where (deoxy(ribose-phosphate linkages are replaced by
N-(2-aminoethy) alycine units linked to nucleotide bases via
glycine amino N through a methylenecarbonyl linker"
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US-09-349-532-1/C

Sequence 1, Application US/09349532

Patent No. 6294650

GENERAL INFORMATION:
APPLICANT: Shay, Jerry W.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Piatyszek, Mieczyslaw A.
APPLICANT: Ocrey, David R.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60

CORRESPONDENCE ADDRESS:
ADDRESSE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STREET: Two Embarcadero Center, Eighth Floor
CONNERS.
CONNERS.
CITY: San Francisco
STREET: Two Embarcadero Center, Eighth Floor
CONNERS.
CONNERS.
CONNERS.
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 2.9%; Score 13; DB 1; Length 13; Best Local Similarity 100.0%; Pred. No. 1.7e+02; Matches 13; Conservative 0; Mismatches 0; Indels
COMPUTER: IBM PC compatible
SOFTWARE: Patentin Release #1.0, Version #1.30
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/838,545
FILING DATE: 09-APR-1997
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/630,019
FILING DATE: 00-APR-1996
ATORNEY AGENT INFORMATION:
NAME: Storella, John R.
REFERENCE/DOCKET NUMBER: 015389-001610US
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 56:
SEQUENCE CHARACTERISTICS:
LEBROTH: 13 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/349,532
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CLASSIFICATION:
PLASSIFICATION DATA:
APPLICATION NUMBER: US 08/838,545
FILING DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE: California
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  55 ACTGAGAAGGGCG 67
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MOLECULE TYPE:
DESCRIPTION:
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DESCRIPTION:
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Length 13;

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Query Match 2.9%; Score 13; DB Best Local Similarity 100.0%; Pred. No. 1.7 Matches 13; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 46, Application US/09349532; Patent No. 6294650; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Floppy disk
                                                                                                                                                                                         44 GICTAACCCIAAC 56
                                                                                                                                                                                                                                42 TTGTCTAACCCTA 54
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CITY: San Francisco
STATE: California
COUNTRY: USA
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TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13; Conservative
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ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            single
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STRANDEDNESS:
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Matches
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                                                                                                                  other nucleic acid (GNA), desc = "peptide nucleic acid (GNA), where (deoxy(ribose-phosphate linkages are replaced by N-(2-aminoethyl) alycine units linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
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where (deoxy(ribose-phosphate linkages are replaced by
N-(2-aminoethyl)glycine units linked to nucleotide bases via
glycine amino N through a methylenecarbonyl linker"
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                                                                                                                                                                                                                                                                                                                                                                                  Query Match 2.9%; Score 13; DB 1; Length 13; Best Local Similarity 100.0%; Pred. No. 1.7e+02; Matches 13; Conservative 0; Mismatches 0; Indels
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ZIP: 050
ZIP: 051
ZIP
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CLASSIFICATION:

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/838,545

FILING DATE: 09-APR-1997

APPLICATION NUMBER: US 08/630,019

FILING DATE: 09-APR-1996

ATTORNEY/AGENT INFORMATION:

NAME: Storella, John R.

REGISTRATION NUMBER: 32,944

REFERENCE/DOCKET NUMBER: 32,944

REGISTRATION INFORMATION:

TELEPHONE: (415) 576-0200

TELEPHONE: (415) 576-0200

INPORMATION FOR EGO ID NO: 12:

SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 12, Application US/09349532
Patent No. 6294650
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                                                       single
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STRANDEDNESS: single
               nucleic acid
TYPE: nucleic acid
STRANDEDNESS: sing
TOPOLOGY: linear
MOLECULE TYPE: ocher
DESCRIPTION: /desc
DESCRIPTION: where
DESCRIPTION: Where
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MOLECULE TYPE:

DESCRIPTION:

DESCRIPTION:

DESCRIPTION:

DESCRIPTION:

US-09-349-532-12
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                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                 APPLICANT: Shay Jerry W. APPLICANT: Shay Jerry W. APPLICANT: Wright, Woodring E. APPLICANT: Piatyszek, Mieczyslaw A. APPLICANT: Corey, David R. APPLICANT: No. 6294650ton, James C. TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids NUMBER OF SEQUENCES: 60 CORRESPONDENCE ADDRESS: ADDRESSE: ADDRESSE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Bighth Floor
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                                                   Indels
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DB 1; Le 1.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/349,532
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:

PRIOR APPLICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/838,545
FILING DATE: 09-ABR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,94
RECTSTATION NUMBER: 32,94
REFERENCE/DOCKET NUMBER: 015389-001610US
TELEFORMUNICATION INFORMATION:
TELEFORM: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "phosphorothioate (PS)
PRSCRIPTION: nucleic acid"
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MOLECULE TYPE: other nucleic acid (PNA), DESCRIPTION: /desc = "peptide nucleic acid (PNA), DESCRIPTION: where (deoxy/ribose-phosphate linkages are replaced by DESCRIPTION: where (deoxy/ribose-phosphate linkages are replaced by DESCRIPTION: N-(2-aminoethy/) glycine units linked to nucleotide bases via DESCRIPTION: glycine amino N through a methylenecarbonyl linker"
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APPLICANT: Wright, Woodring E.
APPLICANT: Paty9zek, Mieczysław A.
APPLICANT: Crey, David R.
APPLICANT: Crey, David R.
APPLICANT: No. 6294650ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by
TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: Callifornia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2.9%; Score 13; DB 1; Length 13; 100.0%; Pred. No. 1.7e+02; ive 0; Mismatches 0; Indels
                                                                                                                                              STATE: California
CONNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compartible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                     JORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                             CORRENT AFFILCATION DATA:
CLASSIFICATION NUMBER: US/09/349,532
FILING DATE:
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: SECRELLA, JOHN R.
REGISTRATION NUMBER: 32,944
REPRENCE/DOCKET NUMBER: 32,944
REPRENCE/DOCKET NUMBER: 015389-001610US
TELEPONE: (415) 576-0300
INPORMATION FOR SEQ ID NO: 52:
SEQUENCE CHARACTERISTICS:
LENGTH. 13 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Sequence 55, Application US/09349532
; Patent No. 6294650
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
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Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-349-532-55/c
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TOPOLOGY: linear

MOLECULE TYPE: other nucleic acid (PNA),

DESCRIPTION: /desc = "peptide nucleic acid (PNA),

DESCRIPTION: where (deoxy(ribose-phosphate linkages are replaced by DESCRIPTION: where (deoxy(ribose-phosphate linkages are replaced by DESCRIPTION: N-(2-aminoethyl)glycine units linked to nucleotide bases via DESCRIPTION: glycine amino N through a methylenecarbonyl linker"
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snay, Jerry W.

APPLICANT: Wright, Woodring E.
APPLICANT: Platyazek, Miscayslaw A.
APPLICANT: Platyazek, Miscayslaw A.
APPLICANT: No. 6294650ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by
TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Towneend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
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US-09-349-532-52/c

i Sequence 52, Application US/09349532

i Patent No. 6294650

i GENERAL INFORMATION:

APPLICANT: Shay, Jerry W.

APPLICANT: Platyszek, Mieczyslaw A.

APPLICANT: Corey, David R.

APPLICANT: Corey, David R.

TITLE OF INVENTION: Modulation of Mammalian Telomerase by
TITLE OF INVENTION: Peptide Nucleic Acids

NUMBER OF SEQUENCES: 60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/349,532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PUCIAS PLANTION:
PRICATION NUMBER: US 08/838,545
FILING DATE: 09-ARR-1997
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-ARR-1996
ATTORNEY/AGENT INFORMATION:
NAME: SEVENIA,001 R.
REFERENCE/DOCKET NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELECPHONE: (415) 576-0300
TELEPAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 51:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
TYPE: NUCleic acid
STRANDEDNES: single
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FILING DATE: 09-APR-1996

ATTONNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 56:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 base pairs
TYPE: MUCHEL Caid
                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: other nucleic acid DESCRIPTION: /desc = "peptide nu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 2, Application US/09657445A Patent No. 6608036
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; Sequence 8, Application US/09657445A
; Patent No. 6608036
; GENERAL INFORMATION:
; APPLICANT: Geron Corporation
; APPLICANT: Gryaznov, Sergei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence
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Matches 13; Conservative
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DESCRIPTION:
DESCRIPTION:
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where (deoxy/tibose-phosphate linkages are replaced by
N.(2-aminoethyl) alycine units linked to nucleotide bases via
glycine amino N through a methylenecarbonyl linker"
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Patent No. 6294650

GENERAL INFORMATION:
APPLICANT: Shay, Jerry W.
APPLICANT: Platyszek, Mieczyslaw A.
APPLICANT: Platyszek, Mieczyslaw A.
APPLICANT: No. 6294650ton, James C.
TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES:
CRRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 2.9%; Score 13; DB 1; Length 13; Best Local Similarity 100.0%; Pred. No. 1.7e+02; Matches 13; Conservative 0; Mismatches 0; Indels
                   CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/349,532
FILING DATE:
CLASSIFICATION NUMBER: US/09/349,532
FILING DATE:
PRIOR APPLICATION NUMBER: US 08/838,545
FILING DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELEPAN: (415) 576-0300
INFORMATION FOR SEQ ID NO: 55:
SEQUENCE CHARACTERISTICS:
LENGTH: 13 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTY: USA
ZIP: 94111-3834
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/838,545
FILIND DATE: 09-APR-1997
APPLICATION NUMBER: US 08/630,019
       PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           other nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             53 TAACTGAGAAGGG 65
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       OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE:
DESCRIPTION:
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DESCRIPTION:
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US-09-349-532-56/C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY:
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GENERAL INFORMATION:
APPLICANT: Geron Corporation
APPLICANT: Geron Corporation
APPLICANT: Grazor, Kristina
APPLICANT: Matray, Tracey
TITLE OF INVENTION: Oligonuclectide N3'-P5' Thiophosphoramidates: Their Synthesis and
TITLE OF INVENTION: Oligonuclectide N3'-P5' Thiophosphoramidates: Their Synthesis and
FILE REFERENCE: 039/003
CURRENT APPLICATION NUMBER: US/09/657,445A
CURRENT FILING DATE: 1999-09-10
PRIOR PELICATION NUMBER: US 60/160,444
PRIOR PELICATION NUMBER: US 60/160,444
PRIOR PELICATION NUMBER: US 60/160,444
SHOR FILING DATE: 1999-10-19
NUMBER OF SEQ ID NOS: 9
SOFTWARE: Patentin version 3.1
SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; OTHER INFORMATION: Synthetic oligonucleotide with potential inhibition activity US-09-657-445A-2
/desc = "peptide nucleic acid (PNA),
where (deoxy(ribose-phosphate linkages are replaced by
N-(2-aminoethyl)glycine units linked to nucleotide bases via
glycine amino N through a methylenecarbonyl linker"
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100.0%; Pred. No. 1.7e+02;
iive 0; Mismatches 0;
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APPLICANT: Gryaznov, Sergei
APPLICANT: Pongracz, Krisztina
APPLICANT: Pongracz, Krisztina
APPLICANT: Matray, Tracey
TITLE OF INVENTION: Oligonucleotide N3'-->P5' Thiophosphoramidates: Their Synthesis ar
TILE REFERENCE: 039/004C
CURRENT APPLICATION NUMBER: US/10/463,076
CURRENT FILING DATE: 2003-06-17
PRIOR PELING DATE: 2000-09-08
PRIOR PELING DATE: 1999-09-10
PRIOR PELING DATE: 1999-09-10
PRIOR PELING DATE: 1999-10-19
NUMBER OF SEQ ID NOS: 9
SOFTWARE: Patentin version 3.1
SEQ ID NO 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: MENG, XIANG-JIN
APPLICANT: HALBUR, PATRICK G.
APPLICANT: HALBUR, PATRICK G.
APPLICANT: HALBUR, PATRICK G.
APPLICANT: HALBUR, PATRICK G.
APPLICANT: HOW, MELISSA A.
TITLE OF INVENTION: A POLYNUCLEIC ACID ISOLATED FROM A
TITLE OF INVENTION: PORCINE REPRODUCTIVE AND RESPIRATORY SYNDROME VIRUS (PRRSV),
TITLE OF INVENTION: PROPERED FROM OR CONTAINING THE POLYNUCLEIC ACID OR
TITLE OF INVENTION: PROTEIN,
NUMBER OF SEQUENCES: 77
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE: , OTHER INFORMATION: Synthetic oligonucleotide with potential inhibition activity US-10-463-076-8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 2.9%; Score 13; DB 1; Length 13; Best Local Similarity 100.0%; Pred. No. 1.7e+02; Matches 13; Conservative 0; Mismatches 0; Indels
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COMPUTER: 1BM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/301,435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 424
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/131,625
FILING DATE: 05-0CT-1993
ATTORNEY/AGENT INFORMATION:
NAME: Lavalleye, Jean-Paul M.P.
REGISTRATION NUMBER: 31,451
REFERENCE/DOCKET NUMBER: 4625-021-55X CIP
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 34, Application US/08301435; Patent No. 6592873; GENERAL INFORMATION: APPLICANT: PAUL, PREM S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 46 CTAACCCTAACTG 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13 CTAACCCTAACTG 1
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE: Virginia
COUNTRY: U.S.A.
ZIP: 22202
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STATE: Virginia
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| Batent No. 6835826
| GENERAL INFORMATION:
| APPLICANT: Geron Corporation
| APPLICANT: Geron Corporation
| APPLICANT: Matray, Tracey
| TITLE OF INVENTION: Oligonuclectide N3'-->P5' Thiophosphoramidates: Their Synthesis of FILE REFERENCE: 039/004C
| CURRENT APPLICATION NUMBER: US/10/463,076
| CURRENT FILING DATE: 2003-06-17 |
| PRIOR FILING DATE: 2000-09-08 |
| PRIOR FILING DATE: 1999-09-10 |
| PRIOR FILING DATE: 1999-09-10 |
| PRIOR FILING DATE: 1999-09-10 |
| PRIOR FILING DATE: 1999-10-19 |
| NUMBER OF SEQ ID NOS: 9 |
| SOFTWARE: Patentin Version 3.1 |
| SOFTWARE: PATENTING DATE: 1999-10-19 |
   APPLICANT: Pongracz, Krisztina
APPLICANT: Matray, Tracey
TITLE OF INVENTION: Oligonucleotide N3'-P5' Thiophosphoramidates: Their Synthesis and
TITLE REFERENCE: 039/003
CURRENT PILING DATE: 2000-09-09
PRIOR APPLICATION NUMBER: US 60/153,201
PRIOR APPLICATION NUMBER: US 60/160,444
PRIOR PILING DATE: 1999-09-10
PRIOR PRILING DATE: 1999-10-19
NUMBER OF SEQ ID NOS: 9
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: Synthetic oligonucleotide with potential inhibition activity US-09-657-445A-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; OTHER INFORMATION: Synthetic oligonucleotide with potential inhibition activity
US-10-463-076-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.9%; Score 13; DB 1; Length 13;
100.0%; Pred. No. 1.7e+02;
tive 0; Mismatches 0; Indels
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; Sequence 8, Application US/10463076
; Patent No. 6815826
; GENERAL INFORMATION;
; APPLICANT: Geron Corporation
                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 2.9°
Best Local Similarity 100.
Matches 13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO 8
LENGTH: 13
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           2.9%; Score 13; DB 1; Length 16; 100.0%; Pred. No. 2.1e+02;
                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                             APPLICANT: McSwiggen, James A.
APPLICANT: Merlo, Patricia Ann Owens
APPLICANT: Guo, Lining
APPLICANT: Skokut, Thomas A.
APPLICANT: Young, Scott A.
APPLICANT: Folkerts, Otto
APPLICANT: Folkerts, Otto
APPLICANT: Merlo, Donald J.
IITLE OF INVENTION: COMPOSITION AND METHODS FOR
IITLE OF INVENTION: MODULATION OF GENE EXPRESSION
IITLE OF INVENTION: IN PLANTS
IITLE OF INVENTION: IN PLANTS
IITLE OF INVENTION: IN PLANTS
       Query Match 2.9%; Score 13; DB Best Local Similarity 100.0%; Pred. No. 2.1 Matches 13; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPOTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/679,645
FILING DATE: July 12, 1996
CLASSIFICATION BATA:
APPLICATION NUMBER: 60/001,135
FILING DATE: July 13, 1995
APPLICATION NUMBER: 08/300,726
FILING DATE: September 2, 1994
ATTORREY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READBALE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                           APPLICANT: Zwick, Michael G.
APPLICANT: Edington, Brent E.
APPLICANT: McSwiggen, James A.
APPLICANT: Merlo, Patricia Ann Owens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 219/247
RELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                           Sequence 828, Application US/08679645; Patent No. 6350934
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SECURORS: 1263
CORRESPONDENCE ADDRESS:
ADDRESSE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  107 GCTGACTTTCAGCGGG 122
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                                                                                                  268 GGGGCTTCTCCGG 280
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 8
SEQUENCE CHARACTERISTICS:
                                                                                                                                               4 GGGGCTTCTCCGG 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 2.8
Best Local Similarity 62.5
Matches 10; Conservative
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TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: MENG, XIANG-JIN
APPLICANT: MENG, XIANG-JIN
APPLICANT: MENG, XIANG-JIN
APPLICANT: HALBUR, PARTICK G.
APPLICANT: HALBUR, PARTICK G.
APPLICANT: HOROZOV, IGOR
APPLICANT: LUM, MELISSA A.
TITLE OF INVENTION: A POLYMUCLEIC ACID ISOLATED FROM A
TITLE OF INVENTION: A PROTEIN ENCODED BY THE POLYMUCLEIC ACID, A VACCINE
TITLE OF INVENTION: PREPARED FROM OR CONTAINING THE POLYMUCLEIC ACID OR
NUMBER OF SEQUENCES: 77
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                    Query Match 2.9%; Score 13; DB 1; Length 16; Best Local Similarity 100.0%; Pred. No. 2.1e+02; Matches 13; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 22202

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/10904
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
CLASSIFICATION:
PRIOR PAPELCATION:
PRIOR PAPELCATION UNMBER: US 08/131,625
APPLICATION UNMBER: US 08/131,625
ATTORNEY/AGENT INFORMATION:
NAME: Lavalleye, Jean-Paul M.P.
REGISTRATION NUMBER: 31,451
REFERENCE/DOCKET NUMBER: 4625-021-55X CIP
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 34, Application PC/TUS9510904
GENERAL INFORMATION:
              TELEFAX: (703) 413-2220
TELEFAX: 248855 OPAT UR
INPORMATION FOR SEQ ID NO: 34:
SEQUENCE CHARACTERISTICS:
LENGHT: 16 base pairs
TYPE: nucleic acid
STRANDEDNESS: unknown
                                                                                                                                                                            TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
(703) 413-3000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: (703) 413-2200
TELEFAX: (703) 413-2220
TELEX: 248655 OPAT UR
INFORMATION FOR SEQ ID NO: 34:
SEQUENCE CHARACTERISTICS:
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PCT-US95-10904-34
                                                                                                                                                                                                                                                                                                                                                                 268 GGGCTTCTCCGG 280
                                                                                                                                                                                                                                                                                                                                                                                         4 GGGCTTCTCCGG 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nucleic acid
EDNESS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 16 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: U c . . ZID
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TELEPHONE:
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PCT-US95-10904-34
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                                            US-08-770-565-2
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    Sequence 1, Application US/09220510B

Sequence 1, Application US/09220510B

Sequence 1, Application US/09220510B

Sequence 10. 6440726

GENERAL INFORMATION:

APPLICANT: RESNICK, NITZAN

TITLE OF INVENTION: EXPRESSION VECTORS COMPRISING MULTIPLE SHEAR STRESS

TITLE OF INVENTION: TREAPONSIVE ELEMENTS (SSRE) AND METHODS OF USE FOR

TITLE OF INVENTION: ANGIOGENESIS IN A SHEAR STRESS ENVIRONMENT

FILE REFERENCE: P-2771-US

CURRENT APPLICATION NUMBER: US/09/220,510B

CURRENT PILING DATE: 1998-12-24

NUMBER OF SEQ ID NOS: 6

SOSTWARE: PALENTIN Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Wu, Fred
APPLICANT: Wu, Fred
APPLICANT: Kealey, James T.
APPLICANT: Realey, James T.
APPLICANT: Pruzan, Ronald
APPLICANT: Weinrich, Scott L.
TITLE OF INVENTION: Methods for Detecting the RNA Component of
TITLE OF INVENTION: Telomerase
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: TWO EMBARCAGERO Center, 8th Floor
STREET: Two Embarcadero Center, 8th Floor
CITY: San Francisco
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 2.2e+02; Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: Description of Artificial sequence: OTHER INFORMATION: A PDGF-A Shear Stress Response Element.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: USA

ZIP: 94111-3834

CONDUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
COMPUTER: PATEMING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATEMIN Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/08/770,565
FILING DATE: 20-DEC-1996
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-002300US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 2, Application US/08770565
Patent No. 5846723
GENERAL INFORMATION:
APPLICANT: Kim, Nam Woo
APPLICANT: Wu, Fred
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 333 GGGGCGAGGCCGAGG 348
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 415-576-0300
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-220-510B-1
US-09-220-510B-1
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US-08-770-565-2
                                                                                                                                                                                                                                                                                                                                 SEQ ID NO 1
LENGTH: 17
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RESULT 220
32-08-873-709-20
is Sequence 20, Application US/08873709
patent No. 6037126
general information:
APPLICANT: Grossman, Abraham
TITLE OF INVENTION: APPARATUS FOR DETERMINING THE PRESENCE OF TITLE OF INVENTION: PROTEIN COMPONENT OF TELOMERASE ENZYME
TITLE OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSE: Abraham Grossman
ADDRESSE: Abraham Grossman
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                                                                                                                                               Length 30;
                                                                                                                                                                                           Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/873,709 FILING DATE: 12-JUN-1997 CLASSIFICATION: 435
                                                                                                                                               Query Match 2.8%; Score 12.8; DB 1; Best Local Similarity 70.8%; Pred. No. 2.9e+02; Matches 17; Conservative 0; Mismatches 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 12.8; DB 1;
Pred. No. 1.7e+02;
0; Mismatches 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Janiuk, Anthony J.
REGISTRATION NUMBER: 29,809
REFERENCE/DOCKET NUMBER: Q001/002
TELECOMUNICATION INFORMATION:
TELECHONE: 914.747-9108
INFORMATION FOR SEQ ID NO: 20:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                222 TCGCCTGCCCAGCCCCGAACCCC 245
                                                                                                                                                                                                                                     131 CCTCGGCCTGCCGCCTTCCACCGT 154
                                                                                                                                                                                                                                                              38 rcccaggcccacccrccgcaaccc 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 16, Application US/09255464B; Patent No. 623867; GENERAL INFORMATION; APPLICANT: Roninson, Igor APPLICANT: Grossman, Abraham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 666 Washington Avenue
CITY: Pleasantville
STATE: NY
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 2.8%;
Best Local Similarity 70.8%;
Matches 17; Conservative
LENGTH: 30 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLGY: linear
MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 62 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
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MOLECULE TYPE:
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US-09-255-464B-15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPOSITIONS, METHODS, KITS AND
APPARATUS FOR DETERMINING THE PRESENCE OR ABSENCE OF
PROTEIN COMPONENT OF TELOMERASE ENZYME
TITLE OF INVENTION: Compositions, Methods, Kits and Apparatus for TITLE OF INVENTION: Identifying Naturally Occurring RNA Sequences Having TITLE OF INVENTION: Affinity for RNA-Binding Proteins FILE REPERENCE: 0001/004a
CURRENT APPLICATION NUMBER: US/09/255,464B
CURRENT FILING DATE: 1999-02-22
NUMBER OF SEQ ID NOS: 25
SOFTWARE: PATENTIN OF 22
SOFTWARE: PATENTIN OF 22
SOFTWARE: PATENTIN OF 22
SOFTWARE: PATENTIN OF 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                              7; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/873,709
FILING DATE: 12-UN-1997
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                            Query Match
2.8%; Score 12.8; DB 1;
Best Local Similarity 70.8%; Pred. No. 1.7e+02;
Matches 17; Conservative 0; Mismatches 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY AGENT INFORMATION:
NAME: Janiuk, Anthony J.
REGISTRATION NUMBER: 29,809
REFERENCE/DOCKET NUMBER: 0001/002
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
INFORMATION FOR SEQ ID NO: 19:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         222 TCGCCTGCCCAGCCCCCGAACCCC 245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               222 TCGCCTGCCCAGCCCCCGAACCCC 245
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 19, Application US/08873709
Patent No. 6037126
GRENEAL INFORMATION:
APPLICANT: Grossman, Abraham
ITILE OF INVENTION: COMPOSITIONS, MITLE OF INVENTION: APPRARATUS FOR LITLE OF INVENTION: APPRARATUS FOR LITLE OF INVENTION: PROTEIN COMPONE NUMBER OP SEQUENCES: 25
CORRESPONDENCE ADDRESS:
ADDRESSEE: Abraham Grossman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29 TCCCAGGCCCACCCTCCGCAACCC 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SSEE: Abraham Grossman
F: 666 Washington Avenue
Pleasantville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              66 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; TOPOLOGY: linear; MOLECULE TYPE: DNA US-08-873-709-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 222
US-08-873-709-19/c
                                                                                                                                                                                                                                                                                                                                               US-09-255-464B-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: N'
COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET:
                                                                                                                                                                                                                                                                           LENGTH: 62
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REBUIT 234

Sequence 15, Application US/09255464B

Sequence 15, Application US/09255464B

Sequence 15, Application US/09255464B

SEQUENCE 17 GEOGRAPHICA STATE STA
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US-08-630-019A-10/c

| Sequence 10, Application US/08630019A
| Sequence 10, Application US/08630019A
| Patent No. 6015710
| GENERAL INFORMATION:
| APPLICANT: Might, Woodring E. APPLICANT: Platyszek, Meczyelaw A. APPLICANT: Corey, David | APPLICANT: No. 6015710ton, James C. TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids | NUMBER OF SEQUENCES 46 | CORRESPONDENCE ADDRESS: ADDRESSE: Townsend and Townsend and Crew LLP | STREET: Townsend and Townsend and CITY: San Francisco | STREET: Control of STREET: Control o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 12; DB 1; Length 12;
Pred. No. 1.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
                            STATE:

COUNTRY: USA

ZIP 94111-3834

ZIP 94111-3834

ZOMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC Compatible

COMPUTER: IBM PC COMPATIBLE

COMPUTER: PATENTIN PC-DOS/MS-DOS

SOFTWARE: PATENTIN NATA:

APPLICATION NUMBER: US/08/770,565

FILING DATE: 20-DEC-1996

CLASSIFICATION NUMBER: 32,944

REGISTRATION NUMBER: 32,944

REGISTRATION NUMBER: 32,944

REFERENCE/DOCKET NUMBER: 15:36-002300US

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION 14:576-0300

TELEFAX: A15-576-0300

TELEFAX: A15-576-0300
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COMPUTER READABLE FORM:
MEDDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATH: US/08/630,019A
FILING DATE: 09-JUN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001600US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 2.7%; Score 12; DB Best Local Similarity 100.0%; Pred. No. 1.8 Matches 12; Conservative 0; Mismatches
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1781: 536
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12 GGAAGAGGAACG 1
         San Francisco
California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-770-565-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Kealey, James T.
APPLICANT: Pruzan, Ronald
APPLICANT: Weinrich, Scott L.
TITLE OF INVENTION: Methods for Detecting the RNA Component of
TITLE OF INVENTION: Telomerase
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: TOWNSEND and management of
STREET: Two Detecting the RNA Component of
TITLE OF INVENTION: Telomerase
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NS-09-121-174A-38/C

Sequence 38, Application US/09328174A

Sequence 38, Application US/09328174A

Sequence 38, Application US/09328174A

GENERAL INFORMATION:
APPLICANT: Guida, Marco
APPLICANT: Kurth, Janice
TITLE OF INVENTION: Genotyping Human Phenol Sulfotransferase
TITLE OF INVENTION: GENOTYPING (STP2)
FILE REFERENCE: 4389-6 (formerly SEQ-16P)
CURRENT FILING DATE: 1999-06-08
PRIOR APPLICATION NUMBER: 09/328,174

PRIOR RILING DATE: 1999-06-08

NUMBER OF SEQ ID NOS: 110

SOFTWARE: PastSEQ for Windows Version 3.0

SEQ ID NO 38
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Pred. No. 2.1e+02;
0; Mismatches 1; Indels
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FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
NAME: WARDING, MICHARD J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 209/166
TELECOMMUNICATION INFORMATION:
TELEPAN: (213) 489-1600
TELEFAX: (213) 955-0440
TELER: 67-3510
INPORMATION FOR SEQ ID NO: 658: SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TYPE: USOFOLOGY: linear
US-08-311-486C-658
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 4, Application US/08770565
Patent No. 5846723
GENERAL INFORMATION:
APPLICANT: Kim, Nam Woo
APPLICANT: Wu, Fred
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       302 AGAGTTGGGCTCTG 315
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 92.57
Best Local 3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15 AGAGTTGGACTCTG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
ORGANISM: H. sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-328-174A-38
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US-08-770-565-4/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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/desc = "peptide nucleic acid (PNA),
where (deoxy/tibose-phosphate linkages are replaced by
N-(2-aminoethy/) glycine units linked to nucleotide bases via
glycine amino N through a methylenecarbonyl linker"
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0
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Patent No. 6294650
GENERAL INFORMATION:
APPLICANT: Shay, Jerry W.
APPLICANT: Shay, Woodring E.
APPLICANT: Platyszek, Mieczyslaw A.
APPLICANT: No. 6294650ton, James C.
ITILE OF INVENTION: Modulation of Mammalian Telomerase by ITILE OF INVENTION: Peptide Nucleic Acids
ITILE OF INVENTION: Peptide Nucleic Acids
NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
ADDRESSER: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STREET: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .;
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                                                                                Length 12;
                                                                                                                            Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA
ZIP: 9411-3834
COMPUTER: READABLE FORM:
MEDIUM TYPE: Flopy disk
COMPUTER: READABLE FORM:
MEDIUM TYPE: Plopy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/349,532
CLASSIPICATION NUMBER: US 08/838,545
FILING DATE: 09-APR-1996
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
APPLICATION NUMBER: US 08/630,019
FILING DATE: 09-APR-1996
APPLICATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELEPRAN: (415) 576-0300
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 12 base pairs
TYPE: mulleic acid
                                                                     2.7%; Score 12; DB 1; Le
100.0%; Pred. No. 1.8e+02;
iive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   other nucleic acid
                                                                                                                       12; Conservative
                                                                                                                                                                        46 CTAACCCTAACT 57
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                                                                   Query Match
Best Local Similarity
Matches 12; Conserv
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TOPOLOGY: lin
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DESCRIPTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DESCRIPTION:
DESCRIPTION:
US-09-349-532-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DESCRIPTION:
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                       US-08-838-545-8
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Best Local S
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                                                                                                                                                                          other nucleic acid (PNA), desc = "peptide nucleic acid (PNA), where (deoxy)ribose-phosphate linkages are replaced by where (deoxy)ribose-phosphate linkages are replaced by N-(2-aminoethyl) glycine units linked to nucleotide bases via glycine amino nitrogen through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          other nucleic acid (descriptions), desc = "peptide nucleic acid (PNN), where (deoxy(ribose-phosphate linkages are replaced by where (12-aminoethy) alycine units linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Shay, Jerry W. APPLICANT: Shay, Jerry W. APPLICANT: Shay, Jerry W. APPLICANT: Shay, Weddring E. APPLICANT: Corey, David M. Statement Corey, David M. APPLICANT: Corey, David M. APPLICANT: No. 6046307ton, James C. TITLE OF INVENTION: Modulation of Mammalian Telomerase by TITLE OF INVENTION: Peptide Nucleic Acids NUMBER OF SEQUENCES: 60 CORRESPONDENCE ADDRESS: ADDRESSE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                Query Match 2.7%; Score 12; DB 1; Length 12; Best Local Similarity 100.0%; Pred. No. 1.8e+02; Matches 12; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURIATA: U.S.A.

ZIP: 94111-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/838,545

FILING DATE: 09-APR-1997

CLASSIFICATION: 336

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/630,019

FILING DATE: 09-APR-1996
ATTORNEY/AGENT INPORMATION:
NAMR: AFCARAILA -- JOHN R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001610US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 8, Application US/08838545
Patent No. 6046307
GENERAL INFORMATION:
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 10
SEQUENCE CHARACTERISTICS:
LENGTH: 12 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: other nucleic
DESCRIPTION: /desc = "pept
DESCRIPTION: where (deoxy)
DESCRIPTION: where (deoxy)
DESCRIPTION: 01ycine aminoc
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TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
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US-08-838-545-8/c
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                                                                                                                                                                                                                           APPLICANT: Sean Sullivan
APPLICANT: Sean Sullivan
APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: RIBASSES OR CONDITIONS
TITLE OF INVENTION: RELATED TO LEVELS OF
TITLE OF INVENTION: RELATED TO LEVELS OF
TITLE OF INVENTION: INTRACELLULAR ADHESION
TITLE OF INVENTION: MOLECULE-1 (I-CAM-1)
NUMBER OF SEQUENCES: 2390
CORRESPONDENCE ADDRESS:
ADDRESSES: Lyon & Lyon
STREET: Sulte 4700
TREET: Sulte 4700
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Sequence 396, Application US/09071845

Patent No. 6132506/C

GENERAL INFORMATION:
APPLICANT: Susan Grimm
APPLICANT: Dan T. Stinchcomb
APPLICANT: Seam Sullivan
APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: TRIBOZYME TREATMENT OF
TITLE OF INVENTION: DISEASES OR CONDITIONS
TITLE OF INVENTION: RELATED TO LEVELS OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19 CTGGGAGGGTG 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12 crescadesers 1
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    Sequence 396, Application US/08292620A

Sequence 396, Application US/08292620A

Patent No. 5837542

GENERAL INFORMATION:
APPLICANT: Susan Grimm
APPLICANT: James McSwiggen
APPLICANT: James McSwiggen
APPLICANT: James McSwiggen
APPLICANT: James McSwiggen
APPLICANT: Sean Sullivan
APPLICANT: Sean Sullivan
APPLICANT: Sean Sullivan
APPLICANT: James McSwiggen
APPLICANT: James McSwiggen
APPLICANT: James McSwiggen
APPLICANT: Sean Sullivan
APPLICANT: Sean Sullivan
APPLICANT: Sean Sullivan
APPLICANTON: RELATED TO LEVELS OF
TITLE OF INVENTION: RELATED TO LEVELS OF
TITLE OF INVENTION: MOLECULE-1 (I-CAM-1)
NUMBER OF SEQUENCES: 2390
CORRESPONDENCE ADDRESS:
ADDRESSEE: 633 West Fifth Street
STREET: 633 West Fifth Street
STREET: 631 West Fifth Street
STREET: 631 West Fifth Street
STREET: 631 West Form:
APPLICANTON TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: ALOUGH
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM Compatible
OPERATING SYSTEM: IBM Compatible
OPERATING SYSTEM: 130,1994
FILING DATE: August 17, 1994
CLASSIFICATION NUMBER: 07/989,849
FILING DATE: Dacember 7, 1992
APPLICATION NUMBER: 07/989,849
FILING DATE: Manuary 19, 1993
APPLICATION NUMBER: 07/989,849
FILING DATE: Warbura 7, 1992
ATTORNEY WARDING PICALATION
NAME: Warbura 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 12; DB 1; Le
Pred. No. 2.3e+02;
0; Mismatches 0;
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REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET WUMBER: 208/149
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 455-0440
TELEFAX: (213) 955-0440
INFORMATION FOR SEQ ID NO: 396:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-292-620A-591/C
; Sequence 591, Application US/08292620A
; Patent No. 5837542
PATENTAL INFORMATION:
APPLICANT: Susan Grimm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 2.7%; Scc
Best Local Similarity 100.0%; P?
Matches 12; Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12 cresedesecre 1
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Gaps
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Best Local Similarity 100.0%; Pred. No. 2.38+02;
Matches 12; Conservative 0; Mismatches 0; Indels
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                                                                                                 COMPUTER READBLE PORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: storage COMPUTER: 1 Storage COMPUTER: IBM COMPATIBLE OPERATING SYSTEM: IBM P.C. DOS 5.0 SOFTWARE: WORD PETECT 5.1 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/071,845 FILING DATE: AUGUST 17,1994
APPLICATION NUMBER: US/08/292,620 FILING DATE: August 17,1994
APPLICATION NUMBER: 08/08,895 FILING DATE: December 7,1993 APPLICATION NUMBER: 07/989,849 FILING DATE: December 7,1993 APPLICATION NUMBER: 07/989,849 FILING DATE: December 7,1992 APPLICATION NUMBER: 07/989,849 FILING DATE: December 7,1992 APPLICATION NUMBER: 22,327 APPLICATION NUMBER: 32,327 APPLICATION N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19 CTGGGAGGGTG 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12 crescadedere 1
                                              California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear
                                                                       N: U.S.A. 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 234
US-09-081-646-720/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-071-845-591
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                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       g
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 233
US-09-071-845-591/C
Sequence 591, Application US/09071845
Fatent No. 612367
GENERAL INFORMATION:
APPLICANT: Susan Grimm
APPLICANT: James McSwigen
APPLICANT: James McSwigen
APPLICANT: Asen Sullivan
APPLICANT: Renneth G. Draper
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: DISEASES OR CONDITIONS
TITLE OF INVENTION: RIATED TO LEVELS OF
TITLE OF INVENTION: INTRACELLULAR ADHESION
TITLE OF INVENTION: NOLECULE: (I-CAM-1)
NUMBER OF SEQUENCES: 2390
CORRESOURNEE ADDRESS:
TITLE OF INVENTION: INTRACELLULAR ADHESION TITLE OF INVENTION: MOLECULE-1 (I-CAM-1) NUMBER OF SEQUENCES: 2390 CORRESPONDENCE ADDRESS: ADDRESSE: Lyon & Lyon STREET: 633 West Fifth Street STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: storage
COMPUTER: IBM COMPALIBLE
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/071,845
FILING DATE:
CLASSIFICATION DATA:
APPLICATION NUMBER: US/08/292,620
FILING DATE: ANGUST 17, 1994
APPLICATION NUMBER: 08/008,895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 208/149
FILENG DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
CRISTRATION NUMBER: 208/149
TELEPHONE: (213) 989-1600
TELEFAK: (213) 985-0440
INFORMATION FOR SEQ ID NO: 396:
SEQUIENCE CHARACTERISTICS:
                                                                                                                                                                                           CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E: Lyon & Lyon
633 West Fifth Street
Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 100.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19 CTGGGAGGGTG 30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; TOPOLOGY:
US-09-071-845-396
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
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2.7%; Score 12; DB 1; Length 16; 91.7%; Pred. No. 2.4e+02; tive 1; Mismatches 0; Indels
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Sequence 4, Application US/08310501

Sequence 4, Application US/08310501

SEQUENCE INFORMATION:

APPLICANT: Sessier, Jonathan L.

APPLICANT: Sessier, Jonathan L.

APPLICANT: Wright, Meredith

STREET: P.O. Box 4433
                                       COMPOSITION AND METHODS FOR MODULATION OF GENE EXPRESSION IN PLANTS
                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: BECAGE
COMPUTER: BECAGE
COMPUTER: BECAGE
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/679,645
FILING DATE: July 12, 1996
CLASSIFICATION DATA:
APPLICATION NUMBER: 06/001,135
FILING DATE: SEPTEMBER: 08/300,726
FILING DATE: SEPTEMBER: 08/300,726
FILING DATE: SEPTEMBER: 08/300,726
FILING DATE: SEPTEMBER: 2, 1994
ATTORNEY/AGENT INFORMATION:
NAME: WATDUREY, RICHARD J.
RECISTRATION NUMBER: 32,327
RECISTRATION NUMBER: 32,327
RECISTRATION NUMBER: 32,327
TELECHONE: (213) 489-1600
                                                                                                                                                                                                                                                                               COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage.
APPLICANT: FOLKETS, OCTO
APPLICANT: Merlo, Donald J.
TITLE OF INVENTION: COMPOSITION AN
TITLE OF INVENTION: MODULATION OF
TITLE OF INVENTION: IN PLANTS
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (213) 955-044(
TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           134 CGGCCTGCCGCC 145
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 91.7
Matches 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 16 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                     CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear
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CITY: Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; TOPOLOGY:
US-08-679-645-515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-310-501-4/c
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                                                                                                                                                                                 Sequence 112, Application US/08050073

Sequence 112, Application US/08050073

GENERAL INFORMATION:
APPLICANT: Begovich, Ann B.
APPLICANT: Begovich, Ann B.
APPLICANT: Brich, Henry A.
APPLICANT: Griffith, Robert L.
APPLICANT: Griffith, Robert L.
APPLICANT: Griffith, Robert L.
APPLICANT: Stephen J.
TITLE OF INVENTION: Typing
NUMBER OF SEQUENCES: 315
CORRESPONDENCE ADDRESS:
ADDRESSEE: Hoffmann-La Roche Inc.
STREET: 340 Kingsland Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP. 77110
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/NS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION NATA:
APPLICATION NUMBER: US/08/050,073
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Zwick, Michael G.
Edington, Brent E.
McSwiggen, James A.
Merlo, Patricia Ann Owens
Guo, Lining
Skokut, Thomas A.
Young, Scott A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 515, Application US/08679645
; Patent No. 6350934
; GENERAL INFORMATION:
; APPLICANT: Zwick, Michael G.
APPLICANT: Edington, Brent E.
; APPLICANT: Merlo, Patricia Ann Owens
; APPLICANT: Skokut, Thomas A.
; APPLICANT: Skokut, Thomas A.
; APPLICANT: Skokut, Thomas A.
; APPLICANT: Young, Scott A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
CLASSIFICATION: 435
ATTORNEY, AGENT INPORMATION:
NAME: PELTY, DOUGLAS A.
REGISTRATION NUMBER: 35,321
REFERENCE/DOCKET NUMBER: 8769
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Pre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      , MOLECULE TYPE: genomic DNA US-08-050-073-112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 100.
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              430 CCAGGACTCGGC 441
                                                   12 GGCTCACATG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: singl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Nutley
STATE: New Jersey
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     linear
                                                                                                                                                                  US-08-050-073-112/c
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US-08-679-645-515
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US-08-484-551-1/c
; Sequence 1, Application US/08484551
; Patent No. 5714328
; APPLICANT: Magda, Darren
; APPLICANT: Reseler, Jonathan L.;
; TILLE OF INVENTION: RNA PHOTOCLEAVAGE USING TEXAPHYRINS
; NUMBER OF SEQUENCES: 8
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Arnold, White & Durkee
; STREET: P.O. Box 4433
; CITY: Houston
; STATE: Texas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 2.6%; Score 11.8; DB 1; Length 15; Best Local Similarity 86.7%; Pred. No. 2.3e+02; Matches 13; Conservative 0; Mismatches 2; Indels
                COUNTRY: Ounited States of America
COUNTRY: United States of America
TIP: 94086
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: PLEADABLE FORM:
COMPUTER: PLEADABLE FORM:
COMPUTER: PLEADABLE FORM:
COMPUTER: PLEADABLE FORM:
COMPUTER: PREADABLE FORM:
COMPUTER: PLEADABLE FORM:
COMPUTER: PLEATION BATA:
APPLICATION NUMBER: US/08/469,177
FILING DATE: 06-JUN-1995
CLASSIFICATION: DATA:
NAME: LATSON, Jacqueline S.
RETERRATION: Jacqueline S.
REGISTRATION NUMBER: PHAY:057
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: mucleic acid
TYPE: mucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTY: United States of America
ZIP: 77210
ZIP: 77210
COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/484,551
FILING DATE: CONCULTENTY herewith
CLASSIPICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Parker, David L.
REGISTRATION NUMBER: 32,165
REFERENCE/DOCKET NUMBER: PHAY:047/PAR
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "RNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            436 CTCGGCTCACATG 450
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(512) 747-7577
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15 ccceecrcacagare 1
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-469-177-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 239
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 g
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2.6%; Score 11.8; DB 1; Length 15;
Best Local Similarity 86.7%; Pred. No. 2.3e+02;
Matches 13; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: SESSLER, Jonathan L.
APPLICANT: SESSLER, Jonathan L.
APPLICANT: IVERSON, Brent L.
APPLICANT: WRIGHT, Meredith
TITLE OF INVENTION: DNA PHOTOCLEAVAGE USING TEXAPHYRINS
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
STREET S
COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
SOCTWARE: Patentin Release #1.0, Version #1.25
SOCTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/08/310,501
FILING DATE: Concurrently herewith
CLASSIFICATION NUMBER: US 08/112,872
FILING DATE: 25-4UG-1993
APPLICATION NUMBER: PCT/US94/06284
FILING DATE: 21-JAN-1992
APPLICATION NUMBER: US 08/227,370
FILING DATE: 14-APR-1994
APPLICATION NUMBER: US 08/075,123
FILING DATE: 14-APR-1992
APPLICATION NUMBER: US 08/771,393
FILING DATE: 30-GEP-1991
APPLICATION NUMBER: US 07/771,393
FILING DATE: 30-SEP-1991
APPLICATION NUMBER: US 07/731,393
FILING DATE: 18-JUN-1990
APPLICATION NUMBER: US 07/539,975
FILING DATE: BUNN-1990
APPLICATION NUMBER: US 07/320,293
FILING DATE: US 07/320,293
FILING DATE: US 07/320,293
FILING DATE: US 07/320,293
FILING DATE: US 07/320,293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PHAY: 034/PAR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 06-MAR-1989
ATTORNEY/AGENT INFORMATION:
NAME: Parker, David L.
REGISTRATION NUMBER: 32,165
REFERENCE/DOCKET NUMBER: PHAY
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 238
US-08-469-177-4/c
; Sequence 4, Application US/08469177
; Patent No. 5607924
; PATENT INFORMATION:
APPLICANT: MAGDA, Darren
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear MOLECULE TYPE: RNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            436 CTCGGCTCACACATG 450
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INFORMATION FOR SEQ ID NO: 6
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: 512/418-3000
TELEFAX: 512/474-7577
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15 ccceecrcacaeare 1
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COMPUTER:
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                                                                                                                                                                                                                            Query Match 2.6%; Score 11.8; DB 1; Length 15; Best Local Similarity 86.7%; Pred. No. 2.3e+02; Matches 13; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 5, Application US/08484551
Patent No. 5714328
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Megda, Darren
APPLICANT: Asseler, Jonathan L.
TITLE OF INVENTION: RNA PHOTOCLEAVAGE USING TEXAPHYRINS
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Arnold, White & Durkee
STREET: P.O. Box 4433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER 1/21D

COMPUTER EXADABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

COMPUTER: TBM PC compatible

COMPUTER: TBM PC compatible

COMPUTER: PAPLICATION DATA:

APPLICATION NUMBER: US/08/484.551

FILING DATE: COMCUTERILY herewith

ATCHASTFICATION: 514

ATCHASTFICATION: 32,165

REGISTRATION NUMBER: 32,165

REGISTRATION NUMBER: 32,165

REGISTRATION NUMBER: 32,165

REGISTRATION NUMBER: 32,165

RELEFAX: (512) 418-3000

TELEFAX: (512) 418-3000

TELEFAX: (512) 418-3000

TELEFAX: (512) 417-7577

TELEY: 79-0924

INFORMATION FOR SEQ ID NO: 5: SEQUENCE CHARACTERISTICS:

LENGTH: 15 base pairs

TYPE: MUCIAL acid

TYPE: MUCIAL acid

TYPE: MUCIAL acid

TYPE: MUCIAL acid
; SEQUENCE CHARACTERISTICS:
; LENGTH: 15 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: other nucleic acid
; DESCRIPTION: /desc = "DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Houston
STATE: Texas
COUNTRY: United States of America
ZIP: 77210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "RNA"
US-08-484-551-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Sequence 18, Application US/08486962
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                                                                                                                                                                                                                                                                                                                                                                          15 cccecrcacacare 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
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US-08-486-962-18/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-484-551-5/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
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RECOMBINANT NEGATIVE STRAND RNA VIRUS
EXPRESSION SYSTEMS AND VACCINES
                                                                                                                                                                                                                                                                                                                            METHOD OF PHOSPHATE ESTER HYDROLYSIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:

MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: Jabenell Batentlu Ba
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STREF: 1155 Avenue of the Americas
CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 11, Application US/08323192D
Patent No. 5786199
GENERAL INFORMATION:
APPLICANT: Palese, Peter
TITLE OF INVENTION: EXPRESSION SYSTEI
NUMBER OF SEQUENCES:
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pharmacyclics, Inc.
                             Sessler, Jonathan L.
Wright, Meredith
Ross, Kevin L.
                                                                                                      APPLICANT: ROSS, Kevin L.
APPLICANT: Miler, Richard A.
APPLICANT: District A.
APPLICANT: Smith, Uadimir A.
APPLICANT: Smith, Daniel A.
TITLE OF INVENTION: METHOD OF P.
UNDRER OF SEQUENCES: 18
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pharmacyclics, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 995 E. Arques Avenue
CITY: Sunnyvale
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              E: Floppy disk
IBM PC compatible
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Magda, Darren
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Best Local Similarity 86.7
Matches 13, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 10036-2711
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA
ZIP: 94086-4521
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE:
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Gaps
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0
                                                                                                                                                          Query Match 2.6%; Score 11.8; DB 1; Length 15; Best Local Similarity 86.7%; Pred. No. 2.3e+02; Matches 13; Conservative 0; Mismatches 2; Indels
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2.6%; Score 11.8; DB 1; Length 15;
Best Local Similarity 66.7%; Pred. No. 2.3e+02;
Matches 10; Conservative 3; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITAIN: SULLEY 4700
CITY: LOS Angeles
STATE: California
COUNTY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/311,486C
FILING DATE: September 23, 1994
CLASSIFTCATION DATA:
PRIOR APPLICATION NUMBER: 08/008,895
FILING DATE: December 7, 1993
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 180. Application US/08311486C
Fatent No. 5811300
GENERAL INFORMATION:
FAPPLICANT: Sean Sullivan
APPLICANT: Kenneth Draper
APPLICANT: Kevin Kisich
TAPLICANT: Town Kisich
APPLICANT: James McSwiggen
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: RIBOZYME TO LEVELS OF
TITLE OF INVENTION: TNF-'
CORRESPONDENCE ADDRESS:
ADDRESSED LYON & LYON
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: LOS Angeles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERNCE/DOCKET NUMBER: 209/166
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                          373 GAGGAACGGAGCGAG 387
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                    TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                 RESULT 244
US-08-311-486C-218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE:
TELEFAX: (2
TELEX: 67-3
                                                                                     ; TOPOLOGY:
US-08-294-424-35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ;
TOPOLOGY:
US-08-311-486C-218
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 35, Application US/08294424
Parent No. 5800984
Parent No. 5800984
Parent No. 5800984
Parent No. 5800984

TILLE OF INVENTION: NUCLEIC ACID SEQUENCE DETECTION BY TITLE OF INVENTION: TRIPLE HELIX FORMATION
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
STREET: 225 Franklin Street
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/323,192D
FILING DATE: 14-OCT-1994
CLASSIFICATION: 435
ATCARSTFICATION: A35
ATCARSTFICATION: LAURA NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 7682-035
TELECOMPUNICATION INFORMATION:
TELEBERONE: (212) 790-9090
TELEBERONE: (212) 790-9090
TELEFAX: (212) 786-9741/8864
TELEX: (212) PRONIE
INFORMATION FOR SEQ ID NO: 11: SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDENSS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: U.S.A.
ZIP: 02110-2804
COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
COMPUTER: 1BM PS/2 Model 50Z or 55SX
OPERATING SYSTEM: 1BM P.C. DOS (Version 3.30)
SOFTWARE: WordPerfect (Version 5.0)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/294,424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/000,922
FILING DATE: 16 JAN 1993
APPLICATION NUMBER: US/07/629,601B
FILING DATE: 17-DEC-1990
ATTORNEY/AGENT INFORMATION:
NAME: Freeman, John W.
REGISTRATION NUMBER: 29,066
REFERENCE/DOCKET NUMBER: 00088-037001
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION: (617) 542-5070
TELEPRIOR: (617) 542-6906
TELEPRIOR: (617) 542-6906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEX: 200154
INFORMATION FOR SEQ ID NO: 35:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         72 CGCCGTGCTTTTGCT 86
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Best Local Similarity 46.74
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Boston
STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: unknown; MOLECULE TYPE: RNA US-08-323-192D-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-08-294-424-35/c
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US-08-316-439A-8

Sequence 8, Application US/08316439A

Patent No. 5840520

GENERAL INFORMATION:

APPLICANT: CLARKE, DAVID KIRKWOOD

APPLICANT: PERER M

TITLE OF INVENTION: SYSTEMS

NUMBER OF SEQUENCES: 43

CORRESPONDENCE ADDRESS.

ADDRESSEE: COOLEY GODWARD CASTRO HUDDLESON & TATUM

STREET: FIVE PALO ALTO SQUARE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IEEE SUCTIONS

COMPUTER: IBM COMPATIBLE

SOFTWARE: Word Perfect 5.1

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/292,620A

FILING DATE: August 17, 1994

CLASSIFICATION DATA:
FRIOR APPLICATION DATA:
FRIOR APPLICATION DATA:
FRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: 08/008,895

FILING DATE: January 9, 1993
APPLICATION NUMBER: 07/989,849

FILING DATE: January 9, 1993
APPLICATION NUMBER: 32,327

REGISTRATION NUMBER: 32,327

REFERENCE/DOCKET NUMBER: 32,327

REGISTRATION NUMBER: 32,327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PatentIn Release #1.0, Version #1.25
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                                                                                                                                                                              CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: SLORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: FIVE PALO ALTO SQUARE
CITY: PALO ALTO
STATE: CALIFORNIA
COUNTRY: USA
ZIP: 94306
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC Compatible
COMPUTER: PROPOSINE: PC-DOS/MS-DOS
SOFTWARE: PATENT Release #1.0, V
                                                                       ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            292:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15 GGGTGGGAGGTGGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                   CORRESPONDENCE ADDRESS:
UMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-292-620A-292
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GENERAL INFORMATION:
APPLICANT: Palese, Peter
APPLICANT: Palese, Peter
APPLICANT: Garcia-saerre, Adolfo
TITLE OF INVENTION: RECOMBINANT NEGATIVE STRAND RNA VIRUS
TITLE OF INVENTION: EXPRESSION SYSTEMS AND VACCINES
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
CITY: New York
COUNTRY: USA
ZIP: 10036-2711
COMPUTER: BW PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CUMBENT APPLICATION DATA:
APPLICATION NUMBER: US/08/470,887A
FILING DATE: 06-UN-1995
CLASSIFICATION
NUMBER: 00-UN-1995
CLASSIFICATION NUMBER: 30,742
REPERENCE/DOCKET NUMBER: 30,742
REPERENCE/DOCKET NUMBER: 30,742
REPERENCE/OCKET NUMBER: 30,742
REPEREN
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Sequence 292, Application US/08292620A

Patent No. 5837542

GENERAL INFORMATION:
APPLICANT: Susan Grimm
APPLICANT: James McSwiggen
APPLICANT: Sean Sullivan
APPLICANT: Sean Sullivan
APPLICANT: Renneth G. Draper
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: RIBOZYME OF LITLE OF INVENTION: RELATED TO LEVELS OF
TITLE OF INVENTION: NUTRACELLULAR ADHESTON
TITLE OF INVENTION: MOLECULE-1 (I-CAM-1)
                                                                                                                                                                                                                                                             US-08-470-887A-10; Sequence 10, Application US/08470887A; Patent No. 5820871
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: RNA (genomic)
                                       436 CTCGGCTCACACATG 450
                                                                               1 CUUGGCUCAGACAUG 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  72 CGCCGTGCTTTTGCT 86
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1 CACCCUGCUUUUGCU 15
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STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 7; Conserva
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46.7%; Pred. No. 2.3e+02;
tive 6; Mismatches 2; Indels
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| Patent No. 5861244
| GENERAL INFORMATION:
| APPLICANT: WANG, C. -G.
| APPLICANT: HEPBIRN, A. G.
| TITLE OF INVENTION: GENERIC SEQUENCE ASSAY USING DNA TITLE OF INVENTION: GENERIC SEQUENCE ASSAY USING DNA TITLE OF INVENTION: TRIPLE-STRAND FORMATION.
| UNMBER OF SEQUENCES: 365
| ORRESPONDENCE ADDRESS: ADDRESSE: ADDRESSE: ROOFLLE DIAGNOSTIC SCIENCES, INC., STREET: 510 EAST 73RD STREET,
| CITY: NEW YORK STATE: NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
MOLECULE TYPE: genomic DNA
DESCRIPTION: hepatitis B virus adr isolate,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CALF: 10021.

COMPUTER: BADABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44Mb storage
COMPUTER: TBM FOXTYAT
OPERATING SYSTEM: MS-DOS version 6.2
SOFTWARE: Wordperfect Version 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/173,489C
FILING DATE: 22 DEC 1993
CLASSIFICATION NUMBER: US/08/173,489C
FILING DATE: 29 DEC 1993
APPLICATION NUMBER: US/08/436
FILING DATE: 29 OCT 1992
ATTORNEY/AGENT INFORMATION:
NAME: Handelman, Joseph H.
REGISTRATION NUMBER: 26,179
REFERENCE/DOCKET NUMBER: U9518-6
TELECOMMUNICATION INFORMATION:
TELECOM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ey) (212) 708-1880
(212) 246-8959
0: 141:
                                                                             FILING DATE: 01-UN-1994
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: COLUZZI, LAURA A.
REGISTRATION NUMBER: 30,742
REPERENCE/DOCKET NUMBER: 7662-034
TELEPHONE: (212) 790-9090
TELEPHONE: (212) 869-9741/8864
TELERAX: (512) 869-9741/8864
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDENESS: single
                                           JMBER: US/08/252,508B
01-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: double stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: unknown

MOLECULE TYPE: RNA (genomic)
US-08-252-508B-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         72 CGCCGTGCTTTTGCT 86
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 46.74
Matches 7; Conservative
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                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 249
US-08-173-489C-141/c
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US-08-252-508B-10

Sequence 10, Application US/08252508B
Fatent No. 5854037
GENERAL INFORMATION:
APPLICANT: Balese, Peter
APPLICANT: Garcia-Gastre, Adolfo
TITLE OF INVENTION: RECOMBINANT NEGATIVE STRAND RNA VIRUS
TITLE OF INVENTION: EXPRESSION SYSTEMS AND VACCINES
NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
CITY: New York
STREET: 1155 Avenue of the Americas
CITY: New York
STREE: Poppy disk
COUNTRY: USA
INDIUM TYPE: Floppy disk
COMPUTER: EMB PC compatible
COMPUTER: IM PC compatible
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.6%; Score 11.8; DB 1; Length 15; 46.7%; Pred. No. 2.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2; Indels
CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/316,439A
FILING DATE: September 30, 1994
CLASSIFICATION 1424

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/190,678
FILING DATE: February 1, 1994
CLASSIFICATION NUMBER: 07/925,061
FILING DATE: August 4, 1992
CLASSIFICATION NUMBER: 07/527,237
FILING DATE: May 22, 1990
CLASSIFICATION NUMBER: 07/527,237
FILING DATE: May 22, 1990
CLASSIFICATION NUMBER: 07/527,237
FILING DATE: May 22, 1990
CLASSIFICATION NUMBER: 07/440,053
FILING DATE: No. 5840520ember 21, 1989
CLASSIFICATION NUMBER: 07/399,728
FILING DATE: August 28, 1989
CLASSIFICATION: 424
PRIOR APPLICATION: 424
PRIOR APPLICATION NUMBER: 07/399,728
FILING DATE: August 28, 1989
CLASSIFICATION NUMBER: AVIN-010/00US
FILING DATE: August 28, 1989
CLASSIFICATION: 424
ATTORNEY, AGENT INFORMATION:
FELENCOMMUNICATION NUMBER: AVIR-010/00US
TELESA: 380816 COOLEY PA
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
TELEX: 380GUSTERISTICS:
TELEX: 580GUSTERISTICS:
TELEX: 580GUSTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6; Mismatches
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MOLECULE TYPE: SYNTHETIC DNA
US-08-316-439A-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               72 CGCCGTGCTTTTGCT 86
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Best Local Similarity 46.73
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 15 bases
TYPE: nucleic acid
STRANDEDNESS: single
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Gaps

Сарв

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RESULT 251
US-09-106-377-10

Sequence 10, Application US/09106377
SEQUENCES 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
STREET: 1155 Avenue of the Americas
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/09/106,377
5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
2.6%; Score 11.8; DB 1;
Best Local Similarity 46.7%; Pred. No. 2.3e+02;
Matches 7; Conservative 6; Mismatches 2;
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Patent No. 6132967
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Susan Grimm
APPLICANT: Dan T. Stinchcomb
APPLICANT: James McSwiggen
APPLICANT: Sean Sullivan
APPLICANT: Kenneth G. Draper
ITLE OF INVENTION: RIBOZYME TREATMENT OF
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/252,508
FILING DATE: 01-010-1994
ATTORNEY/AGENT INFORMATION:
NAME: Coruzzi, Laura A.
REGISTATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 7682-034
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
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INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: RNA (genomic) US-09-106-377-10
                                                      389 CCCCGCGCGCGCGC 403
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       72 CGCCGTGCTTTTGCT 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 CACCCUGCUUUUGCU 15
13; Conservative
                                                                                            1 000000000000000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ð
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US-68-550-120-3
Sequence 3, Application US/08550120
Patent No. 598554
GENERAL INFORMATION:
TITLE OF INVENTION:
MUMBER OF SEQUENCES:
ADDRESSEE: Wenderoth, Lind & Ponack
STREET: 805 Fifteenth Street, N.W., #700
CITY: Washington
STATE: D.C.
COUNTRY: U.S.A.
ZIP: 20005
CONFORTED DISKETCE, S.A.
ZIP: Machington
STATE: D.C.
COUNTRY: U.S.A.
ZIP: Machington
STATE: D.C.
COUNTRY: U.S.A.
ZIP: Machington
STATE: D.C.
COUNTRY: U.S.A.
ZIP: MACHINGTON
MEDIUM TYPE: Diskette, 3.5 inch, 1.44 mb
COMPUTER: IBM Compatible
OPERATING SYSTEM: MS-DOS
SOFTWARE: Wordperfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/550,120
FILING DATE: October 30, 1995
CLASSIFICATION NUMBER: US/08/550,120
FILING DATE: No. 598554ember 2, 1994
ATTORNEY, AGENT INFORMATION:
MAME: Warren M. Cheek, Jr.
REGISTRATION NUMBER: 33,367
REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION NUMBER: 33,367
REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION NUMBER: 33,367
REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION NUMBER: 33,367
REFERENCE/DOCKET NUMBER: 33,367
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                 ORGANISM: Hepatitis B virus
INDIVIDIAL ISOLATE: adr
PUBLICATION INFORMATION:
AUTHORS: Pujiyama, A, Miyanchara, A, No. 5861244aki, C,
AUTHORS: Posyama, T, Ohromo, N, Matsubara, K.
TITLE: Cloning and structural
TITLE: analysis of Hepatitis B virus DNAs subtype adr
VOLUME: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.6%; Score 11.8; DB 1; Length 15; 86.7%; Pred. No. 2.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2; Indels
                                                                                                                                                                                                                                                                                                                                  PAGES: 4601-4610

DATE: 1983

RELEVANT RESIDUES IN SEQ ID NO: 141 :FROM 1 TO 15

US-08-173-489C-141
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.6%; Score 11.8; DB 1;
86.7%; Pred. No. 2.3e+02;
tive 0; Mismatches 2;
  nucleotides 2405 to 2419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 373 GAGGAACGGAGCGAG 387
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15 GAGGAAAGGAGGAG 1
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Best Local Similarity 86.77
The Local Similarity 13. Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
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DESCRIPTION: nu
HYPOTHETICAL: no
ANTI-SENSE: no
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-550-120-3
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OTHER INFORMATION: Description of Artificial Sequence:other nucleic
OTHER INFORMATION: acid
US-09-180-437-212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.6%; Score 11.8; DB 1; Length 15;
86.7%; Pred. No. 2.3e+02;
tive 0; Mismatches 2; Indels
                 COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE 3.5 INCH 1.44 MB STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 212, Application US/09180437
Facent No. 6251873
GENERAL INFORMATION:
APPLICANT: PUKUSANA, Yoshichi
APPLICANT: WORISANA, Yoshichi
TITLE OF INTENTION:
TITLE OF INTENTION: Antisense Compounds to CD14
TITLE OF INTENTION: Antisense Compounds to CD14
TITLE REFERENCE: 1110-209P
CURRENT APPLICATION NUMBER: US/09/180,437
CURRENT FILING DATE: 1998-11-06
EARLIER FILING DATE: 1998-03-09
EARLIER FILING DATE: 1998-03-09
EARLIER FILING DATE: 1998-03-09
SARLIER FILING DATE: 1998-03-09
NUMBER OF SEQ ID NOS: 289
SOFTWARE: Patentin Ver. 2.0
SERVING DATE: 1000-120
                                                          COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: MS-DOS
SOFTWARE: WORDPERFECT 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/871,732A
                                                                                                                                                                                                                                                                                                                                                                                  TELEFACK: 713-682-5807
TELEFACK: 713-682-5807
INPORMATION FOR SEQ 1D NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 BASE PAIRS
TYPE: NUCLEIC ACID
STRANDEDNESS: SINGLE
TOPOLOGY: LINEAR
MOLECULE TYPE: OTHER NUCLEIC ACID
HYPOTHETICAL: NO
HYPOTHETICAL: NO
US-08-871-732A-9
                                                                                                                                                                         FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION 1735
PATJONEY/AGENT INFORMATION:
NAME: MCGREGOR, MARTIN L.
REGISTRATION NUMBER: 29,329
REFRENCE/DOCKET NUMBER: 1-1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 713-682-1213
TELEFAX: 713-682-5807
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21 GGGAGGGGTGGTGGC 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     264 GCCCGGGGCTTCTCC 278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 GGGTGGGGGGGTGGC 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15 GCCGGGGCTTTGCC 1
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Best Local Similarity 86.73
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 254
US-09-180-437-212/c
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Patent No. 6140074
GENERAL INFORMATION:
APPLICANT: O'BRIEN, TIMOTHY J.
APPLICANT: WANG, YIN
APPLICANT: WANG, YIN
TITLE OF INVENTION: CELLS, VECTORS AND EXPRESSION METHOD FOR PRODUCING THE NOVEL
TITLE OF INVENTION: PROTEIN, ANTIBODIES AND USES
NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .;
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         DISEASES OR CONDITIONS
RELATED TO LEVELS OF
INTRACELLULAR ADHESION
MOLECULE-1 (I-CAM-1)
TITLE OF INVENTION: DISEASES OR CONDITIC
TITLE OF INVENTION: RELATED TO LEVELS OF
TITLE OF INVENTION: INTRACELLULAR ADHEST
TITLE OF INVENTION: MOLECULE-1 (I-CAM-1)
NUMBER OF SEQUENCES: 2390
CORRESPONDENCE ADDRESS:
ADDRESSE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STREET: Glifornia
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/071,845
FILING DATE:
CLASSIFICATION DATA:
APPLICATION NUMBER: US/08/292,620
FILING DATE: AMUGER: 17, 1994
APPLICATION NUMBER: 08/008,895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg Richard J.
REFISERRICE/DOCKET NUMBER: 208/149
TELEPHONE: (213) 489-1600
TELEPRAX: (213) 955-0440
TELERAX: 67-3510
INFORMATION FOR SEQ ID NO: 292:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: MARTIN L. MCGREGOR
STREET: 5380 WEST 34TH STREET, #345
CITY: HOUSTON
STATE: TEXAS
COUNTRY: UNITED STATES OF AMERICA
ZIP: 77092
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21 GGGAGGGGTGGTGGC 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15 GGGTGGGGAGGTGGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-071-845-292
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 253
US-08-871-732A-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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Query Match 2.6%; Score 11.8; DB 1; Length 15; Best Local Similarity 86.7%; Pred. No. 2.3e+02; Matches 13; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent NO. 5166057

APPLICANT: PALESE, PETER; PARVIN, JEFFREY D.; KRYSTAL, MARK;

IITLE OF INVENTION: RECOMBIANT NEGATIVE STRAND RNA VIRUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
2.6%; Score 11.8; DB 1;
Best Local Similarity 46.7%; Pred. No. 2.38+02;
Matches 7; Conservative 6; Mismatches 2;
                                                                                                  ; TOPOLOGY: linear ; SEQUENCE DESCRIPTION: SEQ ID NO: 106: US-09-544-934B-106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EXPRESSION-SYSTEMS

NUMBER OF SEQUENCES: 43

CURRENT APPLICATION DATA:

FILING DATE: 22-MAY-1990

PRICK APPLICATION NUMBER: 440, 053

FILING DATE: 21-NOV-1989

APPLICATION NUMBER: 399,728

FILING DATE: 28-AUG-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Search completed: August 24, 2005, 14:29:01 Job time : 3 secs
INFORMATION FOR SEQ ID NO: 106:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 basepairs
TYPE: nucleic acid basepairs
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                           180 CAGCTGCTGGCCCGT 194
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               72 CGCCGTGCTTTGCT 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 CACCCUGCUUUUGCU 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kaare Lund
Tina Anderson Hollerup
TITLE OF INVENTION: No. 6753421el Process For The Detection of Mycobacteria
NUMBER OF SEQUENCES: 123
CORRESPONDENCE ADDRESS:
ADDRESSEE: FINNEGAN, HENDERSON, FARABOW, GARRETT, & DUNNER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                      Sequence 9, Application US/09346510B
Sequence 9, Application US/09346510B
Setent No. 6281014
GENERAL INFORMATION:
APPLICANT: O'Brien, Timothy J.
APPLICANT: Wang, Yinxiang
TITLE OF INVENTION: SH3-Containing Protein, DNA and Uses Thereof
TITLE REFERENCE: D6221CIP
CURRENT APPLICATION NUMBER: US/09/346,510B
CURRENT FILING DATE: 1999-07-01
PRIOR APPLICATION NUMBER: 08/871,732
PRIOR FILING DATE: 1997-06-09
NUMBER OF SEQ ID NOS: 32
SEQ ID NO 9
LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: nucleotide sequence of clone 17 isolated using the COTHER INFORMATION: CASTING approach US-09-346-510B-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 2.6%; Score 11.8; DB 1; Length 15; Best Local Similarity 86.7%; Pred. No. 2.3e+02; Matches 13; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTER READABLE FORM:

MEDIUM TYPE: FLOPPY disk 3.5 inch
COMPUTER: IBM PC compatible
OPERATING SYSTEM: ASCX:
SOFTWARE: Microsoft Word
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/544,934B
FILING DATE: 07-Apr-2000
PRIOR APPLICATION NUMBER: 60/029,595
FILING DATE: 15-Oct-96
APPLICATION NUMBER: 60/029,595
FILING DATE: 23-Oct-96
APPLICATION NUMBER: 60/045,962
FILING DATE: 08-May-97
APPLICATION NUMBER: 08/943,777
FILING DATE: 3-Oct-97
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Washington
STATE: District of Columbia
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 408-4173
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Patent No. 6753421
GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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US-09-827-395A-630 US-09-740-332-800 US-09-740-332-3130 US-09-792-818-129 US-09-792-818-330	us-sn	us-	-Sn	s s	-sn	SO S	us-	-Sn	-Sn	- 50	ns-	-Sn	-SD	181	SS-SS	ns-	-sn	-Sn	20.0	US-10-138-674-455	US-10-138-674-750	US-10-287-949A-45	-sn	US-10-669-841-623	us-	-SD	rsn	-SD	-Sn	nso OS-	ns-	SD S	- SD	ns-	S E	ns-	-sn	-SD	S S	us-	-SD	SO	135	us-	-SD	-SD	US-09-877-478-101	US-09-848-754A-103	US-09-848-754A-103	-sn	US-09-848-754A-165
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13;
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ZIP: 94111-3834

COUNTRY: USA

ZIP: 94111-3834

COMPUTER READALE FORM:
MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
CLASSIFICATION NUMBER: US/09/057,351
PRIOR APPLICATION NUMBER: US/09/330,123
FILING DATE: 07-JUL-1994
PRIOR APPLICATION NUMBER: US/09/472,802
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INPORMATION:
NAME: SECORELLA, JOHN R.
REGISTRATION NUMBER: 32,944
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 22, Application US/09057351
Patent No. US2010034439A1
SENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junia
APPLICANT: Funk, Walter
APPLICANT: Funk, Walter
TITLE OF INVENTION: Mammalian Telomerase
TITLE OF INVENTION: Mammalian Telomerase
UNUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    42 TIGICTAACCCTAACTGAGAAGGGCGTAGGC 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            6.9%; Score 31; DB 100.0%; Pred. No. 13; iive 0; Mismatches
                              TILE OF INVENTION: INDIDITORIES
CURRENT APPLICATION NUMBER: US/10/811,033
CURRENT FILING DATE: 2004-03-26
CURRENT FILING DATE: 2004-03-26
PRIOR APPLICATION NUMBER: US/10/330,872A
PRIOR FILING DATE: 2002-12-24
PRIOR PELING DATE: 1995-08-04
PRIOR PELING DATE: 1995-08-04
PRIOR PLING DATE: 1995-04-04
PRIOR PLING DATE: 1997-04-04
PRIOR APPLICATION NUMBER: 09/420,056
PRIOR APPLICATION NUMBER: 09/420,056
PRIOR PELING DATE: 1999-10-18
PRIOR PELING DATE: 2000-11-20
NUMBER OF SEQ ID NOS: 11
SOFTWARE: PATENTIN VERSION 3.1
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FELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Inhibitors
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Matches 31; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-057-351-22/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
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| Sequence 1, Application US/10330872
| Publication No US2003018628241
| GENERAL INFORMATION:
| Publication No US2003018628241
| GENERAL INFORMATION:
| APPLICANT: Geron Corporation
| APPLICANT: Miniminity, Scott
| APPLICANT: Atlain of Using Purified Telomerase to Identify Telomerase Activators and APPLICANT: Purzan, Ronald
| TITLE OF INVENTION: Unit of Purified Telomerase to Identify Telomerase Activators and TITLE OF INVENTION: Unit of USING PublicATION NUMBER: US/10/330,872
| CURRENT FILING DATE: 1995-08-04
| PRIOR FILING DATE: 1995-08-04
| PRIOR APPLICATION NUMBER: US/883,377
| PRIOR APPLICATION NUMBER: US/883,377
| PRIOR APPLICATION NUMBER: US/800-056
| PRIOR APPLICATION NUMBER: US/800-056
| PRIOR APPLICATION NUMBER: US/800-056
| PRIOR FILING DATE: 1999-10-18
| PRIOR FILING DATE: 1999-10-18
| PRIOR FILING DATE: 1999-10-18
| PRIOR FILING DATE: 2000-11-20
| NUMBER OF SEQ ID NOS: 11
| SEOTTMARE: PATENTIN VERSION 3.1
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| Publication No. US2005008983A1
| GENERAL INFORMATION:
| APPLICANT: Geron Corporation
| APPLICANT: Atkinson III, Edward
| APPLICANT: Lichtsteiner, Serge
| APPLICANT: Pruzan, Rohald
| TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators and
                                                                                                                                                                                                                                                                                             371 AAGAGGAACGGAGGCGAGTCCCCGCGCGCGCGATTCCCTGAGCTGTGGGACGTGCACC 430
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 6.9%; Score 31; DB 1; Length 31; Best Local Similarity 100.0%; Pred. No. 13; Matches 31; Conservative 0; Mismatches 0; Indels
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 84
LENGTH: 79
                                                                                                                                                                                                                                                                                                                                                                                                  431 CAGGACTCGGCTCACACAT 449
                                                                                                                                                                                                                                                                                                                                                                                                                                       ; ORGANISM: Homo sapiens
US-10-714-195-84
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ORGANISM: Homo sapiens
US-10-330-872-1
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                                                                                 TYPE: DNA
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LENGTH: 31
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APPLICANT: Weinrich, Scott
APPLICANT: Atkinson III, Edward
APPLICANT: Atkinson III, Edward
APPLICANT: Atkinson III, Edward
APPLICANT: Lichtsteiner, Serge
APPLICANT: Lichtsteiner, Serge
APPLICANT: Pruzan, Ronald
TITLE OF INVENTION: Unitied Telomerase to Identify Telomerase Activators and
TITLE OF INVENTION: Unititied Telomerase to Identify Telomerase Activators and
TITLE OF INVENTION: UNMBER: 08/10/330,872
CURRENT FILING DATE: 2002-12-24
PRIOR APPLICATION NUMBER: 08/510,736
PRIOR PILING DATE: 1997-04-04
PRIOR PILING DATE: 1997-04-04
PRIOR FILING DATE: 1999-10-18
PRIOR FILING DATE: 09/117,828
PRIOR FILING DATE: 2000-11-20
NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin Version 3.1
SOFTWARE: Patentin Version 3.1
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SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/359,935
                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 015389-000821US
                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
APPLICATION NUMBER: US 08/42,802
FILING DATE: 07-JUN-1995
ATTORNEY AGENT THORNALION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 6.7%; Score 30; DB Best Local Similarity 100.0%; Pred. No. 16; Matches 30; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                             NAME: Storella, John R. REGISTRATION NUMBER: 32,944
                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
                                                                  FILING DATE: 07-Feb-2003
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 4, Application US/10330872
Publication No. US20030186282A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 30 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Geron Corporation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
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APPLICANT: Carimella, Viewanadham
APPLICANT: Garimella, Viewanadham
TITLE OF INVENTION: Method of Detection by Enhancement of Silver Staining
FILE REFERENCE: 00-1086-A
CURRENT APPLICATION NUMBER: US/09/903,461
CURRENT FILING DATE: 2001-07-11
PRIOR APPLICATION NUMBER: 60/217,782
PRIOR FILING DATE: 2000-07-11
SUMBER OF SEQ ID NOS: 3
SOFTWARE: Microsoft Word 98
                                                                                                                                                                                                                                                                                  Gaps
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16;
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                                                                                                                                                                                                                                                                                  0; Indels
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Funk, Walter
Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                  6.7%; Score 30; DB 100.0%; Pred. No. 16; ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                             77 TGCTTTTGCTCCCGCGCGCGCTGTTTTCTC 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        137 CCTGCCGCCTTCCACCGTTCATTCTAGAGC 166
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TTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6.7%; Score 30;
100.0%; Pred. No.
tive 0; Mismatc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; OTHER INFORMATION: synthetic oligomer
US-09-903-461-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 22, Application US/10359935
Publication No. US20030153076A1
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 2, Application US/09903461
Publication No. US20020034756A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA ORGANISM: Artificial Sequence
    TELEFAX: (415) 576-0300
INFORMATION FOR SEG ID NO: 3
SEQUENCE CHARACTERISTICS:
LENGTH: 30 base pairs
                                                                                                                                                                                                                           Query Match
Best Local Similarity 100.
Matches 30, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                          TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
hes 30; Conserva
                                                                                                                                          ; TOPOLOGY: linear; MOLECULE TYPE: DNA US-09-057-351-22
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LENGTH: 30
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Gaps ö

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APPLICANT: Weinrich, Scott
APPLICANT: Weinrich, Scott
APPLICANT: Weinrich, Scott
APPLICANT: Weinrich, Scott
APPLICANT: Atisson III, Edward
APPLICANT: Vasserot, Alain
APPLICANTON NUMBER: US/10/811,033
CURRENT FILING DATE: 2002-12-2
PRIOR PILING DATE: 1997-04-04
PRIOR PELING DATE: 1997-04-04
PRIOR PILING DATE: 1999-10-18
PRIOR FILING DATE: 1999-10-18
PRIOR FILING DATE: 2000-11-20
NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin version 3.1
SOFTWARE: Patentin version 3.1
LENGTH: ANN
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100.0%; Pred. No. 16;
tive 0; Mismatches (
                                                                                                                                                                                                                                                                                                                                                                                                                                              167 AAACAAAAATGTCAGCTGCTGGCCCGTTC 196
                                                                                                                                                                                                                                                                                                                           6.7%; Score 30; DB 100.0%; Pred. No. 16; Live 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30 AAACAAAAATGTCAGCTGCTGGCCCGTTC 1
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                           PRIOR APPLICATION NUMBER: 09/717,828
PRIOR FILING DATE: 2000-11-20
NUMBER OF SEQ ID NOS: 11
SOFTMARE: PatentIn version 3.1
SEQ ID NO BEST OF SEQ ID NOS ID N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 5, Application US/10811033; Publication No. US20050089883A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 3, Application US/10330872; Publication No. US20030186282A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Geron Corporation
APPLICANT: Weinrich, Scott
APPLICANT: Atkinson III, Edward
APPLICANT: Lichtsteiner, Serge
APPLICANT: Vasserot, Alain
APPLICANT: Pruzan, Ronald
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Best Local Similarity 100.
Matches 30; Conservative
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Best Local Similarity 100.
Matches 30; Conservative
                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Homo sapiens
US-10-811-033-4
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; ORGANISM: Homo sapiens
US-10-811-033-5
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US-10-330-872-5/C

i Sequence 5, Application US/10330872

i Publication Wo. US20030186282A1

i GENERAL INFORMATION:

APPLICANT: Weinrich, Scott

APPLICANT: Weinrich, Scott

APPLICANT: Lichtsteiner, Serge

APPLICANT: Dichtsteiner, Serge

APPLICANT: Dichtsteiner, Serge

APPLICANT: Dichtsteiner, Serge

APPLICANT: Using Purified Telomerase to Identify Telomerase Activators and

ITILE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators and

ITILE OF INVENTION: Unibitors

FILE REFERENCE: 011/006C

CURRENT APPLICATION NUMBER: 08/510,736

PRIOR APPLICATION NUMBER: 08/510,736

PRIOR FILING DATE: 1999-04-04

PRIOR FILING DATE: 1999-01-16

PRIOR FILING DATE: 1999-01-16

PRIOR FILING DATE: 1999-01-16

SEQ ID NOS: 11

SEQ ID NO 5

LENGTH: 30

LENGTH: 30

LENGTH: 30
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100.0%; Pred. No. 16;
tive 0; Mismatches 0; Indels
                                                    ;
0
                                                                                                      167 AAACAAAAATGTCAGCTGCTGGCCCGTTC 196
                  Pred. No. 16;
137 CCTGCCGCCTTCCACCGTTCATTCTAGAGC 166
                                                                                                                                     30 AAACAAAAATGTCAGCTGCTGGCCCGTTC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CCTGCCGCCTTCCACCGTTCATTCTAGAGC 1
                                              30; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
ORGANISM: Homo sapiens
US-10-330-872-5
               Best Local Similarity
Matches 30; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 30; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
CLASSITCATION NUMBER: US/09/057,351
FILING DATE: 07-DUL-1994
PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/30,123
FILING DATE: 07-JUN-1995
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:

NAME: SLOCTELIA, JOHN R.

REFERENCE/POCKET NUMBER: 32,944
REFERENCE/POCKET NUMBER: 015389-000821US
TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION SEQ ID NO: 23:
SEQUIRCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                     ADDRESSEE: Townsend and Townsend and Crew LLP STREEF: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 30, Application US/09057351
Fatent No. US20010034439A1
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Juni
APPLICANT: Feng, Juni
APPLICANT: Fank, Walter
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
                                          Sequence 23, Application US/09057351
Fatent No. US2001003439A1
FAPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junii
APPLICANT: Feng, Junii
APPLICANT: Funk, Walter
APPLICANT: Runk, Walter
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
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APPLICANT: Geron Corporation

APPLICANT: Meinrich, Scott

APPLICANT: Aktinson II, Edward

APPLICANT: Aktinson II, Edward

APPLICANT: Vasserot, Alain

APPLICANT: Vasserot, Alain

APPLICANT: Pruzan, Ronald

TITLE OF INVENTION: Units purified Telomerase to Identify Telomerase Activators and

TITLE OF INVENTION: Inhibitors

FILE REPRENCE: 011/006C

CURRENT PPLICATION NUMBER: US/10/811,033

CURRENT FILING DATE: 2002-12-24

PRIOR PRILING DATE: 2002-12-24

PRIOR PELING DATE: 1995-08-04

PRIOR PELING DATE: 1995-08-04

PRIOR PELING DATE: 1995-10-18

PRIOR FILING DATE: 1999-10-18

PRIOR FILING DATE: 1999-10-18

PRIOR FILING DATE: 1999-10-18

PRIOR FILING DATE: 2000-11-20

NUMBER OF SEQ ID NOS: 11

SEQ ID NO 3

LENGTH: 30
TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators and TITLE OF INVENTION: Inhibitors
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; Pred. No. 23;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        412 GAGCTGTGGGACGTGCACCCAGGACTCGGC 441
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 6.3%; Score 28.4; [Best Local Similarity 96.7%; Pred. No. 23; Matches 29; Conservative 0; Mismatches
                                  TILE REPERENCE: 011/006C
CURRENT APPLICATION NUMBER: US/10/330,872
CURRENT FILING DATE: 2002-12-24
PRIOR APPLICATION NUMBER: 08/510,736
PRIOR PILING DATE: 1995-08-04
PRIOR PILING DATE: 1995-04-04
PRIOR PILING DATE: 1997-04-04
PRIOR PILING DATE: 1999-10-18
PRIOR PILING DATE: 1999-10-18
PRIOR PILING DATE: 1990-10-18
PRIOR PILING DATE: 2000-11-20
NUMBER OF SEQ ID NOS: 11
SOFTWARE: PRECENTIN VERSION 3:1
LENGTH: 30
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Publication No. US20050089883A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
ORGANISM: Homo sapiens
US-10-811-033-3
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Best Local Similarity
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AFFLICATION TO THE STATE OF INVENTION: Methods and Reagents for Regulating TITLE OF INVENTION: Methods and Reagents for Regulating Telomere Length and Telomerase Activity
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COMPUTER READABLE FORM:
MEDIUM TYPER: FLOPPY disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/NS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/895,606
FILING DATE: 29-Jun-2001
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/710,249
FILING DATE: CUNROWN-
APPLICATION NUMBER: 08 60/003,492
FILING DATE: 08-SEP-1995
ATTORNEY/AGENT INFORMATION:
NAME: SLOCELLA, JOHN R.
REJERENTENTON NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001220US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 5.8%; Score 26; DB 1; Length 26; Best Local Similarity 100.0%; Pred. No. 30; Matches 26; Conservative 0; Mismatches 0; Indels
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CORRESPONDENCE ADDRESS:
ADDRESSE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                             CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/895,606
FILING DATE: 29-Jun-2001
CLASSIFICATION: «UNKNOWN»
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/710,249
FILING DATE: «UNKNOWN»
APPLICATION NUMBER: US 60/003,492
FILING DATE: 08-SEP-1995
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-001220US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ) MOLECULE TYPE: DNA SEQUENCE DESCRIPTION: SEQ ID NO: 25: US-09-895-606-25
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Publication No. US20030207404A1
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-895-606-26/c
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Reng, Juniiam H.
Adams, William H.
Adams, Robert R.
TITLE OF INVENTION: Methods and Reagents for Regulating
Telomere Length and Telomerase Activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
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ZIP: 94111-3834
COMPUTER READBALE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
ODERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
FILNG DATE: 08-APR-1994
CLASSIFICATION: 435
PRIOR APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-JUN-1995
ATTONINY/AGERI INFORMATION:
ANAME: ALCORDING TO THE TABLE TABLE TO THE TABLE TABLE TO THE TABLE T
JAKEST: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California COUNTRY: USA
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Publication No. US20030207404A1
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: San Francisco
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                                                                                                                                                                                                                                                                                                                                        94111-3834
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US-09-895-606-25
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Matches 2
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Gaps

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APPLICANT: Cech, Thomas R.
Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
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                                               TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 312
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LENGTH: 26 base pai
  SEQUENCE CHARACTERISTICS
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                                                                                         TOPOLOGY: linear
                                                                                                                                                                                                      Query Match
Best Local Similarity 100.0
Matches 26; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                       US-10-044-692-312/c
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Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin
Andrews, William H.
TITLE OF INVENTION: HUMAN TELOMERASE CATALYTIC SUBUNIT: DIAGNOSTIC AND
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                                                                                                                                                                                                                                                                                                                             Gaps
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MEDIUM TYPE: Floppy disk
COMPUTER: DEC COMPATIBLE
COMPUTER: PROPERTIONS
CLASSIFICATION: CURKNOWN>
                                                                                                                                                                                                                                                                         5.8%; Score 26; DB 1; Length 26;
100.0%; Pred. No. 30;
ive 0; Mismatches 0; Indel8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 335
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: Two Embarcadero Center, 8th Floor
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REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002600US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/912,951
FILING DATE: «Unknown»
APPLICATION NUMBER: US 08/654,050
FILING DATE: 09-MAY-1997
APPLICATION NUMBER: US 08/651,843
FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/46,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: 10-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: United States of America
                                                                                                                                                            TOPOLOGY: linear
MOLECULE TYPE: DNA
SEQUENCE DESCRIPTION: SEQ ID NO: 26:
                                                                                                                                                                                                                                                                                                                                                                       145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                    26 CTTCCACCGTTCATTCTAGAGCAAAC 1
TELEPHONE: (415) 576-0200
TELEPAK: (415) 576-0300
INFORMATION FOR SEQ ID NO: 26:
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 311, Application US/10044692
Publication No. US20030096344A1
GENERAL INFORMATION:
APPLICANT: Cech, Thomas R.
Lingner, Joachim
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TELEFAX: (415) 576-0300
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                                                                                                                                                                                                                                                                                                                          26; Conservative
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                                                                                                                                                                                                                                                                                              Best Local Similarity
Matches 26; Conserv
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DIAGNOSTIC AND
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COMPUTER: IBM PC compatible
CERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LLP
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RADDRESSEE: Townsend and Townsend and Crew STREET: Two Embarcadero Center, 8th Floor CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Apple, Randolph T.
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002600US
                                                                                   DB 1;
30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/912,951
FILING DATE: «Uhrinoun»
APPLICATION NUMBER: US 08/854,050
FILING DATE: 09-MAX-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAX-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/844,419
                                                                                   5.8%; Score 26; DB 100.0%; Pred. No. 30; tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: United States of America ZIP: 94111
COMPUTER READABLE FORM:
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APPLICATION NUMBER: US/10/044,692
FILING DATE: 11-Jan-2002
CLASSIFICATION: «Unknown»
MOLECULE TYPE: DNA
SEQUENCE DESCRIPTION: SEQ ID NO: 311:
US-10-044-692-311
                                                                                                                                                                        45 TCTAACCCTAACTGAGAGGGCGTAG 70
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION:
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Gaps

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Harley, Calvin
Harley, Calvin
Andrews, William H.
TITLE OF INVENTION: HUMAN TELOMERASE CATALYTIC SUBUNIT:
THERAPEUTIC METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 26;
                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, 8th Floor
CITY: San Francisco
STATE: California
COUNTRY: United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Apple, Randolph T.
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002600US
TELECOMMUNICATION INFORMATION:
                                                        DB 1;
30;
                                                    ilarity 100.0%; Pred. No. 30; Conservative 0; Mismatches
 SEQUENCE DESCRIPTION: SEQ ID NO: 311:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; TOPOLOGY: linear; single; MOLECULE TYPE: DNA; SEQUENCE DESCRIPTION: SEQ ID NO: 312: US-10-044-539-312
                                                                                                                               45 TCTAACCCTAACTGAGAAGGGCGTAG 70
                                                                                                                                                               1 rcraacccraacreagaagggggrag 26
                                                                                                                                                                                                                                                 Sequence 312, Application US/10044539; Publication No. US20030100093A1; GENERAL INFORMATION: Cech, Thomas R. Lingner, Joachim; Nakamura, Toru Chapman, Karen B. Morin, Gregg B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (415) 576-0300 INFORMATION FOR SEQ ID NO: 312: SEQUENCE CHARACTERISTICS:
                                                      Query Match
Best Local Similarity
Matches 26; Conserv
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US-10-044-539-311
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Nakamura, Toru
Chapman, Kreen B.
Morin, Greeg B.
Harley, Calvin
Andrews, William H.
TITLE OF INVENTION: HUMAN THEOMERASE CATALYTIC SUBUNIT: DIAGNOSTIC AND
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                                                                                                                                              Gaps
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0
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COUNTY: United States of America
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: BAP PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION DATA:
FILING DATE: 11-Jan-2002
CLASSIFICATION: 435
                                                                                                         Length 26,
                                                                                                                                              0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, 8th Floor CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Apple, Randolph T.
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002600US
                                                                                                         DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: AUDKNOWN-
APPLICATION NUMBER: US 08/854,050
FILING DATE: 09-MAY-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: 01-OCT-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                       5.8%; Score 26; DB 100.0%; Pred. No. 30, cive 0; Mismatches
                TOPOLOGY: linear

MOLECULE TYPE: DNA
SEQUENCE DESCRIPTION: SEQ ID NO: 312:
US-10-044-692-312
                                                                                                                                                                              145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/912,951
                                                                                                                                                                                                   26 CTTCCACCGTTCATTCTAGAGCAAAC 1
                                                                                                                                                                                                                                                                                                  Sequence 311, Application US/10044539
Publication No. US20030100093A1
GENERAL INFORMATION:
APPLICANT: Cech, Thomas R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 335
                                                                                                                        Best Local Similarity 100.
Matches 26; Conservative
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US-10-044-539-311
                                                                                                         Query Match
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DIAGNOSTIC AND

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Nakāmura, Toru
Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin B.
Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Version #1.30
                                                                                                                                                                                                                            ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RABGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000821US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 30:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE Floppy disk
COMPUTER: ENORGY disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ver.
CURRENT APPLICATION NUMBER: US/10/359,935
FILING DATE: 07-Feb-2003
CLASSIFICATION: 435
PRIOR APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
APPLICATION NUMBER: US/09/057,351
FILING DATE: 07-ULL-1994
APPLICATION NUMBER: US/08/22,102
FILING DATE: 07-ULL-1994
APPLICATION NUMBER: US/08/32,102
FILING DATE: 07-ULL-1994
                                                                                         Feng, Junli
Funk, Walter
Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 5.8%; Score 26; DB Best Local Similarity 100.0%; Pred. No. 30; Matches 26; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; SEQUENCE DESCRIPTION: SEQ ID NO: 30:
US-10-359-935-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     45 TCTAACCCTAACTGAGAAGGGCGTAG 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Villeponteau, Bryant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 597, Application US/1025810; Publication No. US2030204069A1; GENERAL INFORMATION: APPLICANT: Cech. Thomas R. Lingner, Joachim
         Sequence 30, Application US/10359935
Publication No. US20030153076A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-10-325-810-597
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         Score 26; DB 1; Length 26; Pred. No. 30;
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100.0%; Pred. No. 30;
                                                       Indels
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COMPUTER: IBM PC_COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION UMBER: US/10/359,935
FILING DATE: 07-Feb-2003
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Storella, John R.
REGIGTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000821US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
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APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-UUL-1994
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-UUN-1995
                                                                                                                                                                                                                                                                                                                                                                                   Andrews, William H. TITLE OF INVENTION: Mammalian Telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Pred. ...
         5.8%; Score 26; DB
llarity 100.0%; Pred. No. 30
Conservative 0; Mismatches
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MOLECULE TYPE: DNA
SEQUENCE DESCRIPTION: SEQ ID NO: 23:
US-10-359-935-23
                                                                                                  145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                         CTTCCACCGTTCATTCTAGAGCAAAC 1
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                                                                                                                                                                                                                                US-10-359-935-23/c
; Sequence 23. Application US/10359935
; Publication No. US20030153076A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION
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LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: 23
                                                                                                                                                                                                                                                                                                                                            Feng, Junli
Funk, Walter
                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26; Conservative
Query Match
Best Local Similarity
Matches 26; Conserva
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Best Local Similarity
Matches 26; Conserva
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US-10-359-935-30
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Length 26; Indels

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Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 633
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/10/325,810
FILING DATE: 20-Dec-2002
CLASSIFICATION: «Unknown»
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 5.8%; Score 26; DB 1; Length 26; Best Local Similarity 100.0%; Pred. No. 30; Matches 26; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                       ADDRESSEE: Townsend and Townsend and Crew LL STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: CUNKnown>
PRIOR APPLICATION DATA: US/09/402,181
FILING DATE: 29-Sep-1997
APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-CCT-1996
APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-CCT-1996
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/911,312
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 11-AUG-1997
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 10-OCT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Ausenhus, Scott L.
REGISTRATION NUMBER: 42,271
REFERENCE/DOCKET NUMBER: 015389-002620US
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US-10-325-810-598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                 Nakamura, Toru
Chapman, Karen B.
                                   Lingner, Joachim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (415) 576-0300
                                                                                                                Morin, Gregg B.
             APPLICANT: Cech, Thomas R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 26 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 598
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: -
LOCATION: 1..26
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                      STATE: California COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: DNA
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0
                                                                                                                                                                            ZIP: 9411-3834

COMPUTER READABLE FORM:
MEDLUM TYPE: Floppy disk
COMPUTER: 1BM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRING APPLICATION DATA:
APPLICATION NOWBER: US/10/325,810
FILING DATE: 20-Dec-2002
CLASSIFICATION: CUNKNOWN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 5.8%; Score 26; DB 1; Length 26; Best Local Similarity 100.0%; Pred. No. 30; Matches 26; Conservative 0; Mismatches 0; Indels
NUMBER OF SEQUENCES: 613
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 015389-002620US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE DESCRIPTION: SEQ ID NO: 597:
US-10-325-810-597
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-OCT-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/09/402,181
FILING DATE: 29-SEP-1997
APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-0CT-1996
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/851,843
APPLICATION NUMBER: US 08/851,851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TCTAACCCTAACTGAGAAGGGCGTAG 70
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Ausenhus, Scott L.
REGISTRATION NUMBER: 42,271
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 598, Application US/10325810 Publication No. US20030204069A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      576-0300
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: single
                                                                                                          CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: -
LOCATION: 1..26
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              rypE: nucleic acid
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                                                                                                                                                            COUNTRY: USA
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US-10-325-810-598/c
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Gaps

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RESULT 25
US-10-877-124-597
is Sequence 597, Application US/10877124
is Publication No. US20040242528A1
is GENERAL INFORMATION:
Cech, Thomas R.
ingner, Joachim
Nakamura, Toru
Chapman, Karen B.
Harley, Calvin B.
Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTER READABLE FORM.

EDDING TYPE: L'ALITORILIA

COMPUTER READABLE FORM.

MEDIUM TYPE: L'ADOPY disk
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
COMPUTER: DATE: 24-0Jun-2004
CLASSIPTICATION NUMBER: US/10/877,124
FILING DATE: 24-0Jun-2004
CLASSIPTICATION NUMBER: US/09/432,503
FILING DATE: O2-NOV-1999
APPLICATION NUMBER: US/09/432,503
FILING DATE: O1-NOV-1999
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 16-APR-1997
APPLICATION NUMBER: US 08/911,312
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/911,312
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 11-AUG-1997
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-OCT-1997
APPLICATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 36,429
                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Human Telomerabe cally NUMBER OF SEQUENCES: 727
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLI STREET: Two Embarcadero Center, Eighth Floor CITY: San Prancisco
26 CTTCCACCGTTCATTCTAGAGCAAAC 1
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TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 597
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| Sequence 598, Application US/10877124
| Publication No. US20040242529A1
| GENERAL INFORMATION:
| Lingner, Joachim | Nakamura, Toru | Chapman, Karen B. | Marin, Gregg B. | Harley, Calvin B. | Andrews, William H. | TITLE OF INVENTION: Human Telomerase Catalytic Subunit NUMBER OF SEQUENCES: 727 | CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPLEAS TOWNERS TOWNSEND and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC LODS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/10/877,124
FILING DATE: 24-Jun-2004
FILING DATE: 02-Nov-1999
APPLICATION NUMBER: US/09/432,503
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: CALNROWN-
APPLICATION NUMBER: US 08/844,419
FILING DATE: S-APR-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/845,017
APPLICATION NUMBER: US 08/845,017
APPLICATION NUMBER: US 08/851,843
                                                                                                      Length 26;
                                                                                                                                                     Indele
; SEQUENCE DESCRIPTION: /note= "F3b primer"; SEQUENCE DESCRIPTION: SEQ ID NO: 597: US-10-877-124-597
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: WO PCT/US97/17618
FILING DATE: 01-0CT-1997
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-0CT-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                      DB 1;
30;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
                                                                                                                                                                                                    45 TCTAACCCTAACTGAGAAGGGCGTAG 70
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NAME: Apple, Randolph Ted REGISTRATION NUMBER: 36,429 REFERENCE/DOCKET NUMBER: 015389-002610US

TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 576-0200

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CORRESPONDERS:
ADDRESSE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco COUNTEY: USA
COUNTEY: USA
ZIP: 94111-334
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OOMPUTER: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/877,022
FILING DATE: 24-Jun-2004
CLASSIFICATION *UNMER: US/09/432,503
FILING DATE: 02-Nov-1999
APPLICATION NUMBER: US 09/974,549
FILING DATE: 10-Nov-1999
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APPL-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APPL-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 18-APPL-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 18-APPL-1997
APPLICATION NUMBER: US 08/846,017
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin B.
Andrews, William H.
TITLE OF INVEXTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                          APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-OCT-1997
ATTORNEY/AGENT INFORMATION:
NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002610US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ..
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OTHER INFORMATION: /note= "F3b primer"
SEQUENCE DESCRIPTION: SEQ ID NO: 597:
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30;
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100.0%; Pred. No. ...
... 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TCTAACCCTAACTGAGAAGGGCGTAG 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    45 TCTAACCCTAACTGAGAAGGGCGTAG 70
                                                                                                                                                                                             TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0200
INFORMATION FOR SEG ID NO: 597:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 598, Application US/10877022
Publication No. US20040247613A1
GENERAL INFORMATION:
APPLICANT: Cech, Thomas R.
Lingner, Joachim
            FILING DATE: 01-OCT-1997
                                                                                                                                                                                                                                                                                       LENGTH: 26 base pairs
                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 100.0
Matches 26, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 28
US-10-877-022-598/c
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Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin B.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA
ZIP: 94111-3834
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/877,022
FILING DATE: 24-Jun-2004
CLASSIFICATION: <UNKnown>
                                                                                                                                                                                                                                                                                                                            5.8%; Score 26; DB 1; Length 26;
llarity 100.0%; Pred. No. 30;
Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
                                                                                                                                                                                                                        ; SEQUENCE DESCRIPTION: /note= "R3c primer" 
US-10-877-124-598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: WO PCT/US97/17618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/09/432,503
FILING DATE: 02-Nov-1999
FILING DATE: 08/974,549
FILING DATE: «UNKNOWI)
APPLICATION NUMBER: US 08/44,419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/846,017
RILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 09-MAY-1997
APPLICATION NUMBER: US 08/911,312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 08/854,050
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/912,951
                                                                                                                                                                                                                                                                                                                                                                                                                        145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26 chrocacochroarichaeaceaac 1
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Publication No. US20040247613A1
GENERAL INFORMATION:
INFORMATION FOR SEQ ID NO: 598:
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
                                                                    TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA
FEATURE:
                                                                                                                                                                                   NAME/KEY: -
LOCATION: 1..26
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: California
                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 27
US-10-877-022-597
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Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin B.
Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 727
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1; Length 26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: Ealifornia STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002610US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE DESCRIPTION: SEQ ID NO: 597:
US-10-877-146-597
                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 14-AUG-1997
APPLICATION WUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: WO PCT/US97/17618
FILING DATE: 01-OCT-1997
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-OCT-1997
                                      FILING DATE: 02-Nov-1999
APPLICATION NUMBER: 08/974,549
FILING DATE: «Unknown»
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APF-1997
FILING DATE: 25-APF-1997
                                                                                                                                                                                                   APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-M2-1997
PPLICATION NUMBER: US 08/854,050
FILING DATE: 09-MAY-1997
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/911,312
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/912,951
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5.8%; Score 26; DB
100.0%; Pred. No. 30;
ative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 26 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (415) 576-03
INFORMATION FOR SEQ ID NO: 597
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LOCATION: 1..26
OTHER INFORMATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (415)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26; Conservative
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-877-146-598/c
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Publication No. US20050013825A1
GENERAL INFORMATION:
Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin B.
Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 727
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5.8%; Score 26; DB 1; Length 26;
100.0%; Pred. No. 30;
.ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Townsend and Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco
                                                                                                                                                                                                                                                                                                                NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002610US
TELECOMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 598:
                                    PILING DATE: 09-MAY-1997

APPLICATION NUMBER: US 08/911,312

FILING DATE: 14-AUG-1997

APPLICATION WUMBER: US 08/912,951

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: US 08/915,503

FILING DATE: 14-AUG-1997

APPLICATION NUMBER: WO PCT/US97/17618

FILING DATE: 01-OCT-1997

APPLICATION NUMBER: WO PCT/US97/17618

FILING DATE: 01-OCT-1997

ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LOCATION: 1..26
OTHER INFORMATION: /note= "R3c primer"
SEQUENCE DESCRIPTION: SEQ ID NO: 598:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/10/877,146
FILING DATE: 24-Jun-2004
CLASSIFICATION: «Unknown»
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      145 CTTCCACCGTTCATTCTAGAGCAAAC 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity 100.
Matches 26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 94111-3834
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 29
US-10-877-146-597
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-10-877-022-598
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Indels

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5.5%; Score 25; DB 1; Length 28; 100.0%; Pred. No. 46; 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentun Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-UL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 07-UL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-UTW-1995
ATTORNEY/AGENT INFORMATION:
NAME: SECORIAL, JOHN R.
REFERENCE/DOCKET NUMBER: 32,944
REFERENCE/DOCKET N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Andrews, William H.

TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS: 4
ADDRESSE: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                               NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           81 ITIGCICCCGCGCGCGCTGTTTTCT 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25 rrrigcrecececececrerrrer 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-10-359-935-29/c
; Sequence 29, Application US/10359935
; Publication No. US20030153076A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Feng, Junli
Funk, Walter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                          ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
tes 25; Conserv
                                                                                                                                                                                               STATE: Ca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-057-351-29
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Matches
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0
COMPUTER READABLE FORM:

MEDIUM TYER: RIOPPY disk
COMPUTER: INDEPY disk
SOFTWARE: Patentin Release #1.0, Version #1.30
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/877,146
FILING DATE: 24-Unn-2004
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION DATE:
BPLING DATE: 02-Nov-1999
FILING DATE: 02-Nov-1999
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 12-APR-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-OCT-1997
APPLICATION NUMBER: WO PCT/US97/17885
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 5.8%; Score 26; DB 1; Length 26; Best Local Similarity 100.0%; Pred. No. 30; Matches 26; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002610US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LOCATION: 1..26

COTHER INFORMATION: /note= "R3c primer"

SEQUENCE DESCRIPTION: SEQ ID NO: 598:
US-10-877-146-598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 29, Application US/09057351
Patent No. US20010034439A1
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junii
APPLICANT: Feng, Junii
APPLICANT: Andrews, William H.
TILLE OF INVENTION: Mammalian Telomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             145 CTTCCACCGTTCATTCTAGAGCAAAC 170
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 26 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: 598:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear
MOLECULE TYPE: DNA
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY:
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US-09-057-351-29/c
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0; Gaps

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GENERAL INFORMATION:
APPLICANT: Silverman, Robert H.
APPLICANT: Silverman, Robert H.
APPLICANT: Silverman, Robert H.
APPLICANT: Cowell, John K.
APPLICANT: Torrence, Paul F.
APPLICANT: Torrence, Paul F.
TITLE OF INVENTION: RPRECTIVE TO TREAT TELOMERASE-EXPRESSING MALIGNANCIES;
TITLE OF INVENTION: RPRECTIVE TO TREAT TELOMERASE-EXPRESSING MALIGNANCIES;
TITLE OF INVENTION: BPPECTIVE TO TREAT TELOMERASE-EXPRESSING MALIGNANCIES;
CURRENT APPLICATION NUMBER: US/09/018,125A;
CURRENT FILING DATE: 1999-02-03;
EARLIER APPLICATION NUMBER: 60/044,507;
EARLIER PILING DATE: 1997-04-21
NUMBER OF SEQ ID NOS: 9
SOFTWARE: PATENTING VET. 2.0
SEQ ID NO 5.
LENGTH: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 5.3%; Score 24; DB 1; Length 24; Best Local Similarity 100.0%; Pred. No. 41; Matches 24; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indele
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 345, Application US/10714195
; Sequence 345, Application NS/10714195
; Publication No. US20050019785A1
; GENERAL INFORMATION:
; APPLICANT: Baker, Joffre
; APPLICANT: Cronin, Maureen
; APPLICANT: Cronin, Maureen
; APPLICANT: Gene EXPRESSION PROFILING OF EGFR
; TITLE OF INVENTION: GENE EXPRESSION PROFILING OF EGFR
; TITLE OF INVENTION: DOSITIVE CANCER
; FILE REPERENCE: 39740-0005
; CURRENT APPLICATION NUMBER: US/10/714,195
; CURRENT APPLICATION NUMBER: 60427090
; PRIOR FILING DATE: 2003-11-15
; NUMBER OF SEQ ID NOS: 372
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 345
; LENGTH: 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 5.1%; Score 23; DB 1; Best Local Similarity 100.0%; Pred. No. 48; Matches 23; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        423 CGTGCACCCAGGACTCGGCTCACA 446
TTTGTCTAACCCTAACTGAGAAGG 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         404 GATTCCCTGAGCTGTGGGACGTG 426
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                                                                                                            US-09-018-125-5/c; Sequence 5, Application US/09018125A; Patent No. US20010007902A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA ORGANISM: Artificial Seguence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; OTHER INFORMATION: primer US-10-714-195-345
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 35
US-10-714-195-345/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     g
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Sequence 4, Application US/09018125A

Patent No. U320010007902A1

GENERAL INFORMATION:

APPLICANT: Silverman, Robert H.

APPLICANT: Cowell, John K.

APPLICANT: Cowell, John K.

APPLICANT: Lowell, John K.

CURRENT FOR INVENTION: EFFECTIVE TO TREAT TELOMERASE-EXPRESSING MALIGNANCIES

FILE REFERENCE: 8656-022

CURRENT APPLICATION NUMBER: US/09/018,125A

CURRENT FILING DATE: 1999-02-03

EARLIER FILING DATE: 1997-04-21

NUMBER OF SEQ ID NOS: 9

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 4

LENGTH: 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 5.5%; Score 25; DB 1; Length 28; Best Local Similarity 100.0%; Pred. No. 46; Matches 25; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 5.3%; Score 24; DB 1; Length 24; Best Local Similarity 100.0%; Pred. No. 41; Matches 24; Conservative 0; Mismatches 0; Indels
SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                            PRIOR DATE: 07-Feb-2003
CLASSIFICATION NUMBER: US/10/359,935
FILING DATE: 07-Feb-2003
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APPL-1994
APPLICATION NUMBER: US 08/32,102
FILING DATE: 07-UL-1994
APPLICATION NUMBER: US 08/30,123
FILING DATE: 07-UN-1995
APPLICATION NUMBER: US 08/30,123
FILING DATE: 07-UN-1995
APPLICATION NUMBER: US 08/30,123
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REGISTRATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       , MOLECULE TYPE: DNA
, SEQUENCE DESCRIPTION: SEQ ID NO: 29:
US-10-359-935-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   81 TITGCICCCGCGCGCTGITITCI 105
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (415) 576 0300
INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 28 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-018-125-4
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; OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/siNA sense re
US-10-923-330-511
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WS-10-92-330-512

WS-10-92-330-512

Sequence 312, Application US/10923330

Publication No. US20050153916A1

GENERAL INFORMATION:

APPLICANT: Sirra Therapeutics, Inc.

APPLICANT: McSavigen, James

APPLICANTON: Edgelman, Leonid

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

CURRENT PILING DATE: 2004-08-20

FRIOR PILING DATE: 2004-08-20

PRIOR PELICATION NUMBER: PCT/US03/0408

PRIOR PLING DATE: 2004-05-24

PRIOR PLING DATE: 2004-06-24

PRIOR PLING DATE: 2004-06-24

PRIOR PLING DATE: 2004-06-24

PRIOR PLING DATE: 2004-06-24

PRIOR PLING DATE: 2003-01-14

PRIOR PLING DATE: 2003-11-24

PRIOR PLING DATE: 2003-11-24

PRIOR PLING DATE: 2003-11-23

PRIOR PLING DATE: 2003-11-24

PRIOR PLING DATE: 2003-05-23

PRIOR PLING DATE: 2003-05-23

PRIOR PLING DATE: 2003-05-23

PRIOR PLING DATE: 2003-05-20

PRIOR PLING DA
                      PRIOR FILING DATE: 2004-04-16

PRIOR APPLICATION NUMBER: US 10/757,803

PRIOR APPLICATION NUMBER: US 10/757,803

PRIOR APPLICATION NUMBER: US 10/720,448

PRIOR PELING DATE: 2003-11-24

PRIOR PELING DATE: 2003-11-24

PRIOR PELING DATE: 2003-11-23

PRIOR PELING DATE: 2003-11-23

PRIOR PELING DATE: 2003-11-23

PRIOR PELING DATE: 2003-11-23

PRIOR APPLICATION NUMBER: US 10/44,853

PRIOR APPLICATION NUMBER: PCT/US03/05346

PRIOR PELING DATE: 2003-05-20

PRIOR PILING DATE: 2003-02-20

PRIOR FILING DATE: 2003-02-20

Remaining Prior Application data removed - See File Wrapper or PALM.

NUMBER OF SEQ ID NOS: 768

SOFTWARE: Patentin version 3.3

LENGTH: 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 5.1%; Score 23; DB 1; Length 23; Best Local Similarity 73.9%; Pred. No. 48; Matches 17; Conservative 6; Mismatches 0; Indels
PRIOR APPLICATION NUMBER: US 10/826,966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   136 GCCTGCCGCCTTCCACCGTTCAT 158
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn version 3.3
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LENGTH: 23
TYPE: RNA
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US-10-923-330-510
                                                                             Sequence 510, Application US/10923330

Sequence 510, Application US/10923330

Sequence 510, Application US/10923330

Sequence 510, Application Would US20050153916A1

GENERAL INFORMATION:

APPLICANT: McSwiggen, James

APPLICANTON NUMBER: US/10/23,330

CURRENT FILING DATE: 2004-06-20

PRIOR PAPLICATION NUMBER: US 60/396,600

PRIOR PLILNG DATE: 2004-07-17

PRIOR PLILNG DATE: 2004-01-14

PRIOR PLILNG DATE: 2004-01-14

PRIOR PLILNG DATE: 2003-11-23

PRIOR APPLICATION NUMBER: US 10/720,448

PRIOR PLILNG DATE: 2003-11-23

PRIOR APPLICATION NUMBER: US 10/444,853

PRIOR PLILNG DATE: 2003-11-23

PRIOR APPLICATION NUMBER: US 10/444,853

PRIOR APPLICATION NUMBER: PCT/US03/05346

PRIOR PLILNG DATE: 2003-02-20

PRIOR PLILNG DATE: 2003-02-20

PRIOR PLILNG DATE: 2003-02-20

PRIOR PLILNG DATE: 2003-03-22-30

PRIOR PLING DATE: 2003-03-22-30

PRIOR PLING DATE: 2003-03-22-30

PRIOR PLING DATE: 2003-03-23

PRIOR PLING DATE: 2003-03-23
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Sequence 511, Application US/10923330

Bedication No. US20050153916A1

GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT TILING DATE: 2004-08-20
CURRENT FILING DATE: 2004-08-20

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2005-07-17

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24
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SOFTWARE: PatentIn version 3.3
SEQ ID NO 510
LENCTH: 23
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ORGANISM: Artificial Sequence
             RESULT 36
US-10-923-330-510
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OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/#1NA sense re
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Publication No. US20050153916A1

GENERAL INFORMATION:

APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: McSwiggen, James

APPLICANT: Beigelman, Leonid

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

FILE REFERENCE: 400/209 (MBHB02-708-C)

CURRENT PELING DATE: 2004-08-20

FRIOR PAPLICATION NUMBER: US 60/396,600

FRIOR PELING DATE: 2004-05-24

FRIOR PELICATION NUMBER: US 60/396,600

FRIOR APPLICATION NUMBER: US 60/396,600

FRIOR PELING DATE: 2004-04-16

FRIOR PELICATION NUMBER: US 10/826,966

FRIOR PELICATION NUMBER: US 10/757,803

FRIOR PELICATION NUMBER: US 10/757,803
                                                     ### APPLICANT: Basigelman. Lecond
#### APPLICANT: Basigelman. Lecond
#### TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
#### FILE REFERENCE: 400/209 (MBHB02-708-C)
#### CURRENT APPLICATION NUMBER: US/10/923,330
#### CURRENT FILING DATE: 2004-08-20
### PRIOR PELLACATION NUMBER: US/10/923,330
### CURRENT FILING DATE: 2004-05-24
### PRIOR FILING DATE: 2004-05-24
### PRIOR FILING DATE: 2004-05-24
### PRIOR PELLOCATION NUMBER: US 60/396,600
### RIANG DATE: 2004-05-24
### PRIOR PELLING DATE: 2004-06-24
### PRIOR PELLING DATE: 2004-06-24
### PRIOR PELLING DATE: 2004-01-16
### PRIOR PELLING DATE: 2004-01-16
### PRIOR PELLING DATE: 2004-01-14
### PRIOR PELLING DATE: 2004-01-14
### PRIOR PELLING DATE: 2003-11-24
### PRIOR PELLING DATE: 2003-11-23
### PRIOR PELLING DATE: 2003-11-23
### PRIOR PELLING DATE: 2003-01-23
### PRIOR PELLING DATE: 2003-02-20
### PERLING DATE: 2003-02-20
### PRIOR PELLING DATE: 2
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        McSwiggen, James
Beigelman, Leonid
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                                                                                ; OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/sinA sense US-10-923-330-512
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 513, Application US/10923330
| Publication No. US20050153916A1
| GBNERAL INFORMATION:
| APPLICANT: Sirna Therapeutics, Inc.
| APPLICANT: Sirna Therapeutics, Inc.
| APPLICANT: McSW49GEM, James
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
| FILE REFERENCE: 400/209 (MRH802-708-C)
| CURRENT PILING DATE: 2004-08-24
| FRICH FILING DATE: 2004-08-24
| FRICH FILING DATE: 2004-06-24
| FRICH FILING DATE: 2002-07-17
| FRICH FILING DATE: 2004-04-16
| FRICH FILING DATE: 2004-04-16
| FRICH FILING DATE: 2004-04-16
| FRICH FILING DATE: 2003-04-16
| FRICH FILING DATE: 2003-11-24
| FRICH FILING DATE: 2003-05-20
| FRICH FILING DATE: 2003-05-20
| FRICH FILING DATE: 2003-02-20
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                                                                                                                                                                                                         Query Match 5.1%; Score 23; DB 1; Length 23; Best Local Similarity 95.7%; Pred. No. 48; Matches 22; Conservative 1; Mismatches 0; Indels
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; Sequence 515, Application US/10923330
; Publication No. US20050153916A1
; GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn version 3.3
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; OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/sinA sense re
US-10-923-330-518
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Beiselman, Leonia
1 TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
1 TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
1 TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
1 TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
CURRENT APPLICATION NUMBER: US/10/203, 330
CURRENT PILING DATE: 2004-08-20
FRIOR PELING DATE: 2004-05-24
FRIOR PELING DATE: 2004-05-24
FRIOR PELING DATE: 2004-05-24
FRIOR PELING DATE: 2004-01-17
FRIOR PELING DATE: 2004-01-14
FRIOR PELING DATE: 2004-01-14
FRIOR PELING DATE: 2004-01-14
FRIOR PELING DATE: 2003-01-1-24
FRIOR PELING DATE: 2003-01-1-24
FRIOR PELING DATE: 2003-01-24
FRIOR PELING DATE: 2003-01-3
FRIOR FILING DATE: 2003-01-3
FRIOR FILING DATE: 2003-02-2
FRIOR FILING DATE: 2003-03-2
FRIOR FILING DATE: 2003-03-3
FRIOR FILING DATE:
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leon, Edward Titte of Invention of Telomerase Gene TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REPERENCE: 400,0299 (MBH802-708-C)
CURRENT APPLICATION NUMBER: US,10/923,330
CURRENT FILING DATE: 2004-08-20
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                                       7; Mismatches
                                                                                                      145 CTTCCACCGTTCATTCTAGAGCA 167
                                                                                                                                                         1 CUUCCACCGUUCAUUCUAGAGCA 23
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                                                                                                                                                                                                                                                                                    RESULT 43
US-10-923-330-518
US-10-923-330-518
Sequence 518, Application US/10923330
Publication No. US20050153916A1
GENERAL INFORMATION:
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                                       Conservative
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                               Matches
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'OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/sinA sense

US-10-923-330-516
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US-10-923-330-517
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Sequence 517, Application US/10923330

PUBLIcation No. US20050153916A1

GENERAL INFORMATION:

APPLICANT: Sitra Therapeutics, Inc.

APPLICANT: McSwiggen, James

FILE OF INVENTION: Expression Using Short Interfering RNA (sinA)

FILE REFERENCE: 400/209 (MHHBO: 709-708-7)

FILE REFERENCE: 400/209 (MHHBO: 709-708-7)

FRICK FILING DATE: 2004-08-20

FRIOR FILING DATE: 2004-08-24

FRIOR FILING DATE: 2004-01-14

FRIOR FILING DATE: 2004-01-14

FRIOR FILING DATE: 2004-01-14

FRIOR FILING DATE: 2004-01-14

FRIOR FILING DATE: 2003-11-23

FRIOR FILING DATE: 2003-05-23

FRIOR FILING DATE: 2003-05-34

FRIOR FILING DATE: 2003-05-23

FRIOR FILING DATE: 2003-05-34

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PRIOR FILING DATE: 2003-11-23
PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR PILING DATE: 2003-05-23
PRIOR PILING DATE: 2003-05-20
PRIOR FILING DATE: 2003-02-20
PRIOR FILING DATE: 2003-02-20
PRIOR FILING DATE: 2003-02-20
Remaining Prior Application data removed - See File Wrapper or PALM.
SOFTWARE: Patentin version 3.3
SSQ ID NO 5:16
LENGTH: 23
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48;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                144 CCTTCCACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 42
US-10-923-330-517
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: RNA
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; OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/sinA sense re
US-10-923-330-520
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US-10-923-330-521
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ## APPLICANT: Beagelann. Leonid

## APPLICANT: Beagelann. Leonid

## TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

## TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

## FILE REPERRICE: 400/209 (MHHHD2-708-C)

## CURRENT APPLICATION NUMBER: US/10/923,330

## CURRENT FILING DATE: 2004-08-24

## PRIOR PILING DATE: 2004-05-24

## PRIOR PILING DATE: 2004-05-24

## PRIOR PILING DATE: 2004-05-24

## PRIOR APPLICATION NUMBER: US 60/396,600

## PRIOR PILING DATE: 2004-05-24

## PRIOR PILING DATE: 2004-05-24

## PRIOR PILING DATE: 2004-01-14

## PRIOR PILING DATE: 2004-01-14

## PRIOR PILING DATE: 2004-01-14

## PRIOR PILING DATE: 2003-11-24

## PRIOR PILING DATE: 2003-11-23

## PRIOR PILING DATE: 2003-11-24

## PRIOR PILING DATE: 2003-12-20

## PRIOR PILING DATE: 2003-05-23

## PRIOR PILING DATE: 2003-05-23

## PRIOR PILING DATE: 2003-02-20

## PRIOR PILING DATE: 2003-02
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        Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER OF SEQ ID NOS: 768 SOFTWARE: PatentIn version 3.3 SEQ ID NO 520
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Pred. No. 48;
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48;
                                                                                                                                                                                                                                                                                                                                                                                                                                             5.1%; Score 23; DB 78.3%; Pred. No. 48; tive 5; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  148 CCACCGTTCATTCTAGAGCAAAC 170
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
                                                                                                                                                                                                              TYPE: RNA
ORGANISM: Artificial Sequence
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Best Local Similarity 78.33
Matches 18; Conservative
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Best Local Similarity 78.3
Matches 18; Conservative
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APPLICANT: SIGN INEXPONDED.

APPLICANT: SIGN INEXPONDED.

APPLICANT: Beigelman, Leonid

TITLE OF INVENTION: RNA INFERFERENCE Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

FILE REFERENCE: 400/209 (MBHB02-708-C)

CURRENT PAPLICATION NUMBER: US/10/923,330

CURRENT PAPLICATION NUMBER: DCT/US03/0408

PRIOR APPLICATION NUMBER: DCT/US04/16390

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24

PRIOR PELICATION NUMBER: US 10/826,966

PRIOR PELICATION NUMBER: US 10/826,966

PRIOR APPLICATION NUMBER: US 10/757,803

PRIOR PILING DATE: 2004-01-14

PRIOR FILING DATE: 2003-11-24

PRIOR FILING DATE: 2003-11-23

PRIOR FILING DATE: 2003-11-23

PRIOR FILING DATE: 2003-05-23

PRIOR FILING DATE: 2003-05-23
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PRIOR APPLICATION NUMBER: PCT/USO3/04088
PRIOR PILING DATE: 2004-05-24
PRIOR PLLING DATE: 2004-05-24
PRIOR PLLING DATE: 2002-07-17
PRIOR APPLICATION NUMBER: US 60/396,600
PRIOR PLLING DATE: 2004-05-24
PRIOR PLLING DATE: 2004-05-24
PRIOR APPLICATION NUMBER: US 10/826,966
PRIOR PLLING DATE: 2004-01-14
PRIOR PLLING DATE: 2004-01-14
PRIOR PLLING DATE: 2003-01-23
PRIOR PLLING DATE: 2003-11-24
PRIOR PLLING DATE: 2003-11-23
PRIOR PLLING DATE: 2003-11-23
PRIOR APPLICATION NUMBER: US 10/494,853
PRIOR PLLING DATE: 2003-05-23
PRIOR PLLING DATE: 2003-05-20
PRIOR PLLING DATE: 2003-02-20
PRIOR PLLING DATE: 2003-02-20
REMAINING PLLES 2003-02-20
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APPLICATION NUMBER: PCT/US03/05028
FILING DATE: 2003-02-20
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Publication No. US20050153916A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: RNA
ORGANISM: Artificial Sequence
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LENGTH: 23
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4.9%; Score 22; DB 1; Length 22; 100.0%; Pred. No. 55; 0; Indels ive 0; Mismatches 0; Indels
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco
                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Eloppy disk
COMPUTER: IEM PC compatible
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
   E: Townsend and Townsend and Crew LLP
Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                              | FILING DATE: 08-APR-1994 | CLASSIFICATION UNDER: 08/APR-1994 | CLASSIFICATION: 435 | PRIOR APPLICATION DATA: 435 | PRIOR APPLICATION DATA: 435 | APPLICATION NUMBER: US 08/272,102 | FILING DATE: 07-ULL-1994 | PRIOR APPLICATION DATA: 48-PLICATION DATA: APPLICATION DATA: 48-PLICATION NUMBER: US 08/472,802 | FILING DATE: 07-ULN-1995 | ATTORNEY/AGENT INFORMATION: NAME: Storella, John R. REGISTRATION NUMBER: 32,944 | REGERENCE/DOCKET NUMBER: 015389-000821US | TELEFONMUNICATION INFORMATION: TELEFONMUNICATION NUMBER: 015389-000821US | TELEFONMUNICATION NUMBER: 0
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Funk, Walter
Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 41, Application US/10359935
Publication No. US20030153076A1
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: California
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22 AACTGAGAAGGGCGTAGGCGCC 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 22 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22; Conservative
                        STREET: TWO DAMES CITY: San Francisco CTATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 22; Conserva
ADDRESSEE:
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US-10-359-935-41/c
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100.0%; Pred. No. 55;
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-UN-1995
ATTORNEY/AGENT INFORMATION:
ARGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000821US
TELEPRONE (415) 576-0200
TELEPRONE (415) 576-0200
TELEPRAX: (415) 576-0300
                                          Sequence 1, Application US/09057351
Patent No. US20010034439A1
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Juni;
APPLICANT: Feng, Juni;
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
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US-09-057-351-42/c
; Sequence 42, Application US/09057351
; Parent No. US20010034439A1
; GENERAL INFORMATION:
APPLICANT: Villeporteau, Bryant
APPLICANT: Feng, Junli
; APPLICANT: Funk, Walter
; TITLE OF INVENTION: Mammalian Telomerase
; TITLE OF INVENTION: Mammalian Telomerase
; CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/NS-DOS
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Best Local Similarity
Matches 22; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 USA
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                 US-09-057-351-41/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
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FILING DATE: 07-JUN-1995
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MEDIUM TYPE: Floppy disk
COMPUTER: INPW DC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/359,935
FILING DATE: 07-Feb-2003
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 22;
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TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS: 70
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                         NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000821US
REFERENCE/DOCKET NUMBER: 015389-000821US
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 41:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
55;
APPLICATION NUMBER: US/10/359,935
FILING DATE: 07.Feb-2003
CLASSIPICATION: 435
PRIOR APPLICATION: 435
PRIOR APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-ULL-1994
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-UNN-1995
ATTORNEY, AGENT INFORMATION:
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APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
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APPLICATION NUMBER: US 08/472,802
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear
MOLECULE TYPE: RNA
SEQUENCE DESCRIPTION: SEQ ID NO: 41:
US-10-359-935-41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 4.9%; Score 22; Best Local Similarity 100.0%; Pred. No. Matches 22; Conservative 0; Mismatch
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US-10-359-935-42/C
; Sequence 42, Application US/10359935
; Publication No. US20030153076A1
; GENERAL INFORMATION:
APPLICANT: Fand Junii
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STATE: California
COUNTRY: USA
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Funk, Walter
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GENERAL INFORMATION:
APPLICANT: Geron Corporation
APPLICANT: Geron Corporation
APPLICANT: Geron Corporation
APPLICANT: Maintain, Scott
APPLICANT: Lichtsteiner, Serge
APPLICANT: Display Purified Telomerase to Identify Telomerase Activators and
TITLE OF INVENTION: Uniblicors
FILE REFERENCE: 01/006C
CURRENT APPLICATION NUMBER: US/10/330,872
CURRENT FILING DATE: 1995-08-04
PRIOR PILING DATE: 1995-08-04
PRIOR PRILING DATE: 1999-10-18
PRIOR PRILING DATE: 1999-10-18
PRIOR PLILNG DATE: 2000-11-20
NUMBER OF SEC ID NOS: 11
SEC ID NO 2
LENGTH: 22
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100.0%; Pred. No. 55;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                         Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000821US
TELECOMMUNICATION INFORMATION:
TELEFRAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 42:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
55;
                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Pred. No. 55; tive 0; Mismatches
                                                                                                                                                                                                                                                                           TOPOLOGY: linear
MOLECULE TYPE: RNA
SEQUENCE DESCRIPTION: SEQ ID NO: 42:
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; Publication No. US20050003404A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 2, Application US/10330872; Publication No. US20030186282A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     54 AACTGAGAAGGGCGTAGGCGCC
                                                                                                                                                                                                         LENGTH: 22 base pairs
                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 100.0
Matches 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 100.
Matches 22; Conservative
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; ORGANISM: Homo sapiens
US-10-330-872-2
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Gaps
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55;
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                                        TITLE OF INVENTION: TELOMERASE INTERFERENCE
FILE REPERENCE: A-1156-2/RFT/THR
CURRENT APPLICATION UNMBER: US/10/831,267
CURRENT FILING DATE: 2004-04-22
FRIOR PELIONG NUMBER: US/00/831,267
FRIOR FILING DATE: 2002-10-16
FRIOR FILING DATE: 2001-10-22
FRIOR RILING DATE: 2001-10-22
FRIOR PELICATION NUMBER: US 60/345,326
FRIOR FILING DATE: 2001-0-22
FRIOR APPLICATION NUMBER: US 60/389,195
FRIOR PILING DATE: 2002-02-20
FRIOR APPLICATION NUMBER: US 60/383,195
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Sequence 13, Application US/10831267
Fublication No. US2005000917741
GENERAL INFORMATION:
APPLICANT: ROWLey, Peter T.
TITLE OF INVENTION: TELOMERASE INTERFERENCE
FILE REFERENCE: A-71506-2/RFT/THR
CURRENT APPLICATION NUMBER: US/10/831,267
FRIOR PAPLICATION NUMBER: US/10/831,267
PRIOR PAPLICATION NUMBER: US (0/345,326
PRIOR FILING DATE: 2002-10-16
PRIOR APPLICATION NUMBER: US 60/345,326
PRIOR PILING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: US 60/359,196
PRIOR FILING DATE: 2001-10-22
PRIOR FILING DATE: 2002-05-2
PRIOR FILING DATE: 2002-05-2
NUMBER OF SEQ ID NOS: 23
SOFTWARE: PatentIn Version 3.2
LENGTH: 22
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4.9%; Score 22; DB
Best Local Similarity 100.0%; Pred. No. 55;
Matches 22; Conservative 0; Mismatches
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US-10-811-033-2/c
; Sequence 2, Application US/10811033
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: primer US-10-831-267-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
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US-10-831-267-13/c
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Best Local Si
Matches 22,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Indels
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4.9%; Score 22; DB 1;
Best Local Similarity 100.0%; Pred. No. 55;
Matches 22; Conservative 0; Mismatches (
APPLICANT: Rowley, Peter T.

JITLE OF INVENTION: TELOMERASE INTERFERENCE
FILE REFERENCE: A-71506-1/RFT/THR
CURRENT APPLICATION NUMBER: US/10/831,266
CURRENT FILING DATE: 2004-04-22
RIOR APPLICATION NUMBER: US/10/83365
PRIOR APPLICATION NUMBER: US 60/345,326
PRIOR APPLICATION NUMBER: US 60/359,196
PRIOR FILING DATE: 2001-10-22
PRIOR FILING DATE: 2002-02-20
PRIOR PELLORATION NUMBER: US 60/359,196
PRIOR PLILING DATE: 2002-02-22
PRIOR APPLICATION NUMBER: US 60/383,195
PRIOR RILING DATE: 2002-05-22
NUMBER OF SEQ ID NOS: 17
SOFTWARE: PARCHILIN VERSION 3.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 13. Application US/10831266
Publication No. US2005003404A1
GENERAL INFORMATION:
APPLICANT: ROWLEY. Peter T.
TITLE OF INVENTION: TELOMERASE INTERFERENCE
FILE REFERENCE: A-71506-1/RFT/THR
CURRENT FILING DATE: 2004-04-22
PRIOR APPLICATION NUMBER: US/10/831,266
PRIOR APPLICATION NUMBER: US/0/33065
PRIOR APPLICATION NUMBER: US 60/345,326
PRIOR APPLICATION NUMBER: US 60/345,326
PRIOR FILING DATE: 2002-10-16
PRIOR APPLICATION NUMBER: US 60/359,196
PRIOR FILING DATE: 2002-02-20
PRIOR APPLICATION NUMBER: US 60/359,196
PRIOR FILING DATE: 2002-02-20
PRIOR FILING DATE: 2002-05-22
NUMBER OF SEQ ID NOS: 17
SOFTWARE: PARCENTIN VERSION 3.2
SEQ ID NO 13
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tive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         176 ATGTCAGCTGCTGGCCCGTTCG 197
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
Matches 22; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 53
US-10-831-266-13/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-10-831-266-12
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US-10-831-267-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IN PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT PAPLICATION NATA:
APPLICATION NUMBER: US/10/359,935
FILING DATE: 07-Feb-2003
CLASSIFICATION 435
PRIOR APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APPL1994
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
APPLICATION NUMBER: US 08/30,123
FILING DATE: 27-OCT-1994
APPLICATION NUMBER: US 08/310,123
APPLICATION NUMBER: US 08/310,123
FILING DATE: 07-JUL-1994
APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-JUL-1994
APPLICATION NUMBER: US 08/472,802
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4.7%; Score 21; DB 1; Length 21;
100.0%; Pred. No. 64;
tive 0; Mismatches 0; Indels
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Funk, Walter
Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000821US
                 APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-UN-1995
ATTORNEY/AGENT INFORMATION:
NAMB: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000821US
TELECHANINICATION INFORMATION:
TELEFAX: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 25, Application US/10359935
Publication No. US20030153076A1
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          184 TGCTGGCCCGTTCGCCCCTCC 204
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 100.1
Matches 21; Conservative
                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS LENGTH: 21 base pairs
                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: DNA
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                                      APPLICANT: Geron Corporation
APPLICANT: Weinrich, Scott
APPLICANT: Atkinson III, Edward
APPLICANT: Atkinson III, Edward
APPLICANT: Lichtsteiner, Serge
APPLICANT: Druzan, Ronald
TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators and
TITLE OF INVENTION: Using Purified Telomerase to Identify Telomerase Activators
FILE REFERENCE: 011/006C
CURRENT FILING DATE: 2004-03-26
PRIOR PRIOR APPLICATION NUMBER: 08/10/330,872A
PRIOR PLING DATE: 1995-08-04
PRIOR PILING DATE: 1997-04-04
PRIOR PLING DATE: 1997-04-04
PRIOR FILING DATE: 1997-04-04
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COUNTRY: USA
ZIP: 9411-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
TT.ING DATE: 08-ARR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Jos-09-057-351-25/c

Sequence 25, Application US/09057351

Sequence 25, Application US/09057351

GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junii
APPLICANT: Funk, Walter
APPLICANT: Funk, Walter
TITLE OF INVENTION: Wammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend are
STREET: Two Fermine Application Telomerase
CITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 4.9%; Score 22; DB Best Local Similarity 100.0%; Pred. No. 55; Matches 22; Conservative 0; Mismatches
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              46 CTAACCCTAACTGAGAAGGGCG 67
Publication No. US20050089883A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: Homo sapiens
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Publication No. US20050153916A1
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US-10-923-330-536

Sequence 536, Application US/10923330

Publication No. US2000013916A1

SEGNERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
TITLE OF INVERTION:
TITLE OF INVERTION: Expression Using Short Interfering RNA (sinA)
TITLE OF INVERTION: Expression Using Short Interfering RNA (sinA)
FILE REPERENCE: 400/209 (WHEND2-708-C)
CURRENT PRILING DATE: 2004-06-20
FRIOR APPLICATION NUMBER: PCT/US03/04088
FRIOR PLING DATE: 2002-07-17
FRIOR APPLICATION NUMBER: US 60/396,600
FRIOR FILING DATE: 2000-07-17
FRIOR PLING DATE: 2004-01-4
FRIOR PLING DATE: 2003-11-24
FRIOR APPLICATION NUMBER: US 10/757,803
FRIOR FILING DATE: 2003-11-24
FRIOR PLING DATE: 2003-11-24
FRIOR PLING DATE: 2003-11-24
FRIOR PLING DATE: 2003-12-34
FRIOR FILING DATE: 2003-12-34
FRIOR FILING DATE: 2003-02-30
FRIOR FILING DATE: 2003-02-20
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                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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4.7%; Score 21; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 64;
Matches 21; Conservative 0; Mismatches 0; Indels
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4.7%; Score 21; DB 1; Length 21;
Best Local Similarity 81.0%; Pred. No. 64;
Matches 17; Conservative 4; Mismatches 0; Indels
                                   TYPE: nucleic acid
STRANDEDNESS: single
STRANDEDNESS: ningle
TOPOLOGY: linear
MOLECULE TYPE: DNA
SEQUENCE DESCRIPTION: SEQ ID NO: 25:
                                                                                                                                                                                                                                                                                                                                                                                                                                               184 TGCTGGCCCGTTCGCCCTCC 204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21 recreeccerreccerec 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn version 3.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
                                                                                                                                                                                                                           US-10-359-935-25
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US-10-923-330-537
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; Sequence 537, Application US/10923330

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| ARPLICANT: RESEARCH INTORANTOR, APPLICANT: APPLICANT:
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US-10-923-330-542/c
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Sequence 539, Application US/10923330

Sequence 539, Application US/10923330

Sequence 539, Application US20050153916A1

SPUBLICANT: Sirna Therapeutics, Inc.

APPLICANT: Beigelman, Leonid

TITLE OF INVENTION: RAD Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RAD Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RAD INTERFERENCE: 400/209 (MBHE02-708-C)

CURRENT APPLICATION NUMBER: US/10/23,330

CURRENT FILING DATE: 2004-08-20

PRIOR FILING DATE: 2004-06-20

PRIOR FILING DATE: 2004-05-24

PRIOR APPLICATION NUMBER: US 60/396,600

PRIOR FILING DATE: 2004-05-24

PRIOR APPLICATION NUMBER: US 10/752,903

PRIOR APPLICATION NUMBER: US 10/752,903

PRIOR APPLICATION NUMBER: US 10/752,903

PRIOR FILING DATE: 2003-11-24

PRIOR APPLICATION NUMBER: US 10/752,948

PRIOR APPLICATION NUMBER: US 10/752,948

PRIOR APPLICATION NUMBER: US 10/752,948

PRIOR APPLICATION NUMBER: US 10/753,959

PRIOR APPLICATION NUMBER: US 10/753,959

PRIOR APPLICATION NUMBER: US 10/753,963

PRIOR APPLICATION NUMBER: US 10/753,963
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, OTHER INFORMATION: Description of Artificial Sequence: sinA sense region
US-10-923-330-539
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Description of Artificial Sequence: siNA sense region
Gaps
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
Matches 20; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-10-923-330-538
                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 538
LENGTH: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         a
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GENERAL INFORMATION TO The Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McGaiggan, James
APPLICANT: McGaiggan, James
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REPERENCE: 400/209 (WHRH02-708-C)
CURRENT TLIME DATE: 2004-06-20
FILE REPERENCE: 400/209 (WHRH02-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT CLIME DATE: 2004-06-20
FRIOR PRICK FILING DATE: 2004-07-17
FRIOR FILING DATE: 2004-06-24
FRIOR FILING DATE: 2004-04-16
FRIOR FILING DATE: 2004-05-23
FRIOR FILING DATE: 2004-05-23
FRIOR FILING DATE: 2003-05-24
FRIOR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ó
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Publication No. US20050153916A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Beigelman, Jemes
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (81NA)
                                                                                   Gарв
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Length 21;
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64;
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Score 21; DB 1;
Pred. No. 64;
4; Mismatches (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 4.7%; Score 21; DB Best Local Similarity 100.0%; Pred. No. 64; Matches 21; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 541, Application US/10923330 Publication No. US20050153916A1 GENERAL INFORMATION:
                                                                                                                                                              397 GCGCCCCATTCCCTGAGCTG 417
                                                                                                                                                                                                                 1 GCGGCGCGAUUCCCUGAGCUG 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 GGTTGCGGAGGGTGGGCCTGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
    Query Match
Best Local Similarity 81.0%;
Matches 17; Conservative
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RESULT 66

UR-10-223-330-544/C

Sequence 544, Application US/10923330

Sequence 544, Application US/10923330

Sequence 544, Application US/10923330

GENERAL INFORMATION:
Publication No. US20050153916A1

APPLICANT: SITEM Therapeutics, Inc.
APPLICANT: Beigalman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
FILE EMPERENCE: 400/209 (MRH002-708-C)
CURRENT APPLICATION NUMBER: DC/10/922,330

CURRENT APPLICATION NUMBER: PC/10/03/0408

PRIOR APLICATION NUMBER: PC/10/396,600

PRIOR PILING DATE: 2004-05-24

PRIOR PILING DATE: 2004-05-24

PRIOR PILING DATE: 2004-05-24

PRIOR PILING DATE: 2004-05-24

PRIOR PILING DATE: 2004-05-14

PRIOR PILING DATE: 2004-05-14

PRIOR PILING DATE: 2004-05-12

PRIOR PILING DATE: 2004-05-12

PRIOR PILING DATE: 2004-05-12

PRIOR PILING DATE: 2004-05-14

PRIOR PILING DATE: 2004-05-14

PRIOR PILING DATE: 2004-05-14

PRIOR PILING DATE: 2003-11-23

PRIOR PILING DATE: 2003-01-23

PRIOR PELING DATE: 2003-01-23

PRIOR PELING DATE: 2003-01-23

PRIOR PELING DATE: 2003-02-20

PRIOR APPLICATION NUMBER: US 10/44, 853

PRIOR PELING DATE: 2003-02-20

PRIOR APPLICATION NUMBER: PCT/US03/05346

PRIOR APPLICATION NUMBER: PCT/US03/05346

PRIOR APPLICATION NUMBER: PCT/US03/05246

PRIOR PRIOR PRIOR PRIOR POST 768

SEQ ID NOS: 768

SEQ ID NOS: 768

SEQ ID NOS: 768

SED ID NOS: 768

SED ID NOS: 768

PUBLICATION

PUBLICATION POST 768

PUBLICATION PUBLICATION NUMBER: PCT/US03/05246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region US-10-923-330-543
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
US-10-923-330-544
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                                      PRIOR FILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR FILING DATE: 2003-02-20
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 768
SOFTWARE: Patentin version 3.3
SEQ ID NO 543
LENGTH: 21
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4.7%; Score 21; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 64;
Matches 21; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             283 GCACCCACTGCCACCGCGAAG 303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           395 GCGCGCGCGATTCCCTGAGC 415
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                                                                                                                                                                                                                                                                                                                                                                               TYPE: RNA
ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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Best Local Similarity
Matches 21; Conserval
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Sequence 543, Application US/10923330

Publication No. US20050153916A1

GENERAL INFORMATION:

APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: McSwiggen, James

APPLICANT: Beigelman, Leonid

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

CURRENT PRILICATION NUMBER: DCT/US03/04088

PRIOR APPLICATION NUMBER: PCT/US03/04088

PRIOR APPLICATION NUMBER: DCT/US03/04088

PRIOR FILING DATE: 2004-05-24

PRIOR PRILING DATE: 2004-05-24

PRIOR PRILING DATE: 2004-05-24

PRIOR PRILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-01-16

PRIOR FILING DATE: 2004-01-16

PRIOR FILING DATE: 2004-01-12

PRIOR PRILOR APPLICATION NUMBER: US 10/757,803

PRIOR FILING DATE: 2003-11-24

PRIOR FILING DATE: 2003-11-24

PRIOR APPLICATION NUMBER: US 10/444,853

PRIOR FILING DATE: 2003-305-23

PRIOR FILING DATE: 2003-05-23

PRIOR PRIOR FILING DATE: 2003-05-346
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CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-08-20
FRIOR APPLICATION NUMBER: PCT/US03/04088
FRIOR APPLICATION NUMBER: PCT/US03/04088
FRIOR PRICOR FILING DATE: 2004-05-24
FRIOR PELING DATE: 2004-05-24
FRIOR PELING DATE: 2004-05-24
FRIOR PELING DATE: 2004-05-24
FRIOR PELING DATE: 2004-04-14
FRIOR PELING DATE: 2004-04-14
FRIOR PELING DATE: 2003-10-14
FRIOR FILING DATE: 2003-11-24
FRIOR FILING DATE: 2003-05-3
FRIOR FILING 
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4.7%; Score 21; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 64;
Matches 21; Conservative 0; Mismatches 0; Indels
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 65
US-10-923-330-543/c
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ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California
                                                                                                                                                                                                                                     COUNTRY: USA

ZIP: 94111-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATION SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTION DATA:
APPLICATION NATA:
APPLICATION NATA:
APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/330,123
FILING DATE: 07-JUL-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-JUN-1955
ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 015389-000821US
TELECOMMUNICATION INFORMATION:
TELEBROME: (415) 576-0200
TELEBROME: (415) 576-0300
INFORMATION FOR SEQ ID NO: 40:
SEQUENCE CHARACTERISTICS:
LENGTH: C. D DASE DATE

LENGTH: C. DASE DATE
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
4.4%; Score 20; DB 1;
Best Local Similarity 100.0%; Pred. No. 73;
Matches 20; Conservative 0; Mismatches
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Funk, Walter
Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerage
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 41 TTTGTCTAACCCTAACTGAG 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TTTGTCTAACCCTAACTGAG 1
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-359-935-7/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: USA
ZIP: 9411-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
FILING DATE: 07-JUL-1994
PRIOR APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
APPLICATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000821US
TELEPHONE: (415) 576-0300
TELEPHONE: (415) 576-0300
TELEPHONE: (415) 576-0300
TELEPAT: NON-PATE
SEQUENCE CHARACTERISTICS:
CHARACT
                                                                                                                                          Sequence 7, Application US/09057351
; Sequence 7, Application US/09057351
; Patent No. US/2001003439A1
; GENERAL INFORMATION:
APPLICANT: Funk, Walter
APPLICANT: Funk, Walter
APPLICANT: Funk, Walter
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
ADDRESSE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
COUNTRY: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 40, Application US/09057351
Patent No. US20010034499A1
PAPELICANT: Villeponteau, Bryant
APPLICANT: Feng, Junii
APPLICANT: Feng, Junii
APPLICANT: Peng, William H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Villeponteau, Bryant
Feng, Junli
Funk, Walter
Andrews, William H.
          21 GCGCGCGCGATTCCCTGAGC 1
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Best Local Similarity 100.
Matches 20; Conservative
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STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 68
US-09-057-351-40/c
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Gaps ö

Length 20; Indels

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APPLICATION NUMBER: US 08/330,123
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                                                                                                                                                                                                                      TELEFAX: (415) 576-0300 INFORMATION FOR SEQ ID NO: 40: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  41 TITGICTAACCCTAACTGAG 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20 TTTGTCTAACCCTAACTGAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 100.0%; Pr
Matches 19; Conservative 0;
                                                                                                                                                                                                                                                                                               LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  76 GIGCITITGCICCCGCGC
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Best Local Similarity 100.(
Matches 20; Conservative
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              CURRENT APPLICATION DATA:

APPLICATION DATA:

APPLICATION NUMBER: US/10/359,935

FILING DATE: 07-Feb-2003

CLASSIFICATION: 435

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/09/057,351

FILING DATE: 08-APR-1994

APPLICATION NUMBER: US 08/372,102

FILING DATE: 07-JUL-1994

APPLICATION NUMBER: US 08/330,123

FILING DATE: 27-OCT-1994

APPLICATION NUMBER: US 08/310,123

FILING DATE: 27-OCT-1994

APPLICATION NUMBER: US 08/310,123

FILING DATE: 27-OCT-1994

ATTORNEY/AGENT INFERMATION:

AND ATTORNEY/AGENT INFERMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4.4%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 73; 0; Indels ive 0; Mismatches 0; Indels
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: 1BM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATCHTIN Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                    NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000821US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Feng, Junli
Funk, Walter
Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/057,351
FILING DATE: 08-APR-1994
APPLICATION NUMBER: US 08/272,102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/359,935
FILING DATE: 07-Feb-2003
CLASSIFICATION: 435
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; TOPOLOGY: Linear; MOLECTLE TYPE: DNA SEQUENCE DESCRIPTION: SEQ ID NO: 7: US-10-359-935-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-10.359-935-40/c
US-10.359-935-40/c
; Sequence 40, Application US/10359935
; Publication No. US20030153076A1
; GENERAL INFORMATION:
; APPLICANT: Villeponteau, Bryant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 07-JUL-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GGTTGCGGAGGGTGGGCCTG 21
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SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
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Best Local Similarity
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GENERAL INFORMATION:

APPLICANT: Silverman, Robert H.

APPLICANT: Silverman, Robert H.

APPLICANT: Solid, Seiji

APPLICANT: Cowell, John K.

APPLICANT: Cowell, John K.

APPLICANT: Torrence, Raul F.

TITLE OF INVENTION: RNASE L ACTIVATORS AND ANTISENSE OLIGONUCLEOTIDES

TITLE OF INVENTION: RNASE L (18/09/018,125A)

TITLE OF INVENTION: BFFECTIVE TO TREAT TELOMERASE-EXPRESSING MALIGNANCIES

FILE REFERENCE: 8656-022

CURRENT APPLICATION NUMBER: 08/09/018,125A

CURRENT FILING DATE: 1999-02-03

EARLIER APPLICATION NUMBER: 60/044,507

EARLIER FILING DATE: 1997-04-21

NUMBER OF SEQ ID NOS: 9

SEQ ID NO 2.
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Pred. No. 82;
0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: oligonucleotide
                                                                        ATTORNEY/AGENT INFORMATION:
NAME: Storella, John R.
REGISTRATION NUMBER: 32,944
REFERENCE/DOCKET NUMBER: 015389-000821US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4.4%; Score 20; DB 100.0%; Pred. No. 73; Live 0; Mismatches
FILING DATE: 27-OCT-1994
APPLICATION NUMBER: US 08/472,802
FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
MOLECULE TYPE: RNA
SEQUENCE DESCRIPTION: SEQ ID NO: 40:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 72
US-10-016-490C-8
; Sequence 8, Application US/10016490C
; Publication No. US20040072769AL
; GENERAL INFORMATION:
; APPLICANT: Yin, James Q.
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TYPE: DNA ORGANISM: Artificial
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                                                                                                                                                                                                                             TYPE: RNA
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TITLE OF INVENTION: Methods for design and selection of short double-stranded TITLE OF INVENTION: oligonucleotides, and compounds of gene drugs FILE REFERENCE: 01-279 CURRENT APPLICATION NUMBER: US/10/016,490C CURRENT FILING DATE: 2002-11-22 NUMBER OF SEQ ID NOS: 51 SOFTWARE: Patentin version 3.1
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                                                                                                                                                                                                                                                                                                                                          Length 19;
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                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                        Query Match
4.2%; Score 19; DB 1;
Best Local Similarity 100.0%; Pred. No. 82;
Matches 19; Conservative 0; Mismatches
                                                                                                                                                                                                                                                    ; OTHER INFORMATION: The same as those in human. US-10-016-490C-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NESOLI 5)
Sequence 16, Application US/10831267
Sequence 16, Application US/10831267
Sequence 16, Application US/10831267
GENERAL INFORMATION:
TITLE OF INVENTION: TELOMERASE INTERFERENCE
FILE REFERENCE: A-71506-2/RRT/THR
CURRENT APPLICATION NUMBER: US/10/831,267
CURRENT APPLICATION NUMBER: US/10/831,267
CURRENT APPLICATION NUMBER: US 60/345,326
PRIOR PILING DATE: 2004-04-22
PRIOR PILING DATE: 2004-16
PRIOR FILING DATE: 2001-10
PRIOR FILING DATE: 2001-10
PRIOR APPLICATION NUMBER: US 60/359,196
PRIOR FILING DATE: 2001-0-2
PRIOR APPLICATION NUMBER: US 60/389,195
PRIOR FILING DATE: 2002-0-20
PRIOR FILING DATE: 2002-0-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: PARCH NOSE 23
SEQ ID NO 16
LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; Sequence 17, Application US/10831267
; Publication No. US20050009177A1
; GENERAL INFORMATION:
; APPLICANT: ROWLEY, Peter T.
; TILE OF INVENTION: TELOMERASE INTERFERENCE
; FILE REFERENCE: A-71506-2/RFT/THR
; CURRENT APPLICATION NUMBER: US/10/831,267
; CURRENT FILING DATE: 2004-04-22
; PRIOR APPLICATION NUMBER: PCT/US 02/33146
; PRIOR FILING DATE: 2002-10-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              6; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                  372 AGAGGAACGGAGCGAGTCC 390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 AGAGGAACGGAGCGAGTCC 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          42 TTGTCTAACCCTAACTGAG 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 UUGUCUAACCCUAACUGAG 19
                                                                                                                                                                                                                             ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
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US-10-831-267-17/c
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                                                                                                                                                                                                          TYPE: DNA
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4.2%; Score 19; DB 1; Length 19;
Best Local Similarity 100.0%; Pred. No. 82;
Matches 19; Conservative 0; Mismatches 0; Indels
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; Sequence 18, Application US/10831267
; Publication No. US20050009177A1
; GENERAL INFORMATION:
    APPLICANT: Rowley, Peter T.
    TITLE OF INVENTION: TELOMERASE INTERFERENCE
    TITLE OF INVENTION: TELOMERASE INTERFERENCE
    CURRENT PAPLICATION NUMBER: US/10/831,267
; CURRENT FILING DATE: 2004-04-22
    PRIOR PELICATION NUMBER: PCT/US 02/33146
; PRIOR APPLICATION NUMBER: DCT/US 02/33146
; PRIOR PELING DATE: 2001-10-22
; PRIOR FILING DATE: 2001-10-22
; PRIOR FILING DATE: 2002-20
; PRIOR FILING DATE: 2002-05-2
; PRIOR FILING DATE: 2002-05-2
; NUMBER OF SEQ ID NOS: 23
; SOFTUMARE: Patentin Version 3.2
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Publication No. US2005009177A1
GENERAL INFORMATION:
TITLE OF INVENTION: TELOMERASE INTERFERENCE
FILE REFERENCE: A.71506-2/RFT/THR
CURRENT FILING DATE: 2004-04-22
PRIOR APPLICATION NUMBER: US/10/831,267
PRIOR APPLICATION NUMBER: PCT/US 02/33146
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4.2%; Score 19; DB
Best Local Similarity 100.0%; Pred. No. 92;
Matches 19; Conservative 0; Mismatches
PRIOR APPLICATION NUMBER: US 60/345,326
PRIOR FILING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: US 60/359,196
PRIOR FILING DATE: 2002-02-20
PRIOR FILING DATE: 2002-05-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: PatentIn version 3.2
SEQ ID NO 17
LENGTH: 19
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Publication No. US20050019785A1
GENERAL INFORMATION:
APPLICANT: Baker, Joffre
APPLICANT: Cronin, Maureen
APPLICANT: Cronin, Maureen
APPLICANT: Shaker, Jose
TITLE OF INVENTION: GENE EXPRESSION PROFILING OF EGFR
TITLE OF INVENTION: GENE EXPRESSION PROFILING
FILE REFERENCE: 39740-0005
CURRENT APPLICATION NUMBER: US/10/714,195
CURRENT PILING DATE: 2003-11-14
PRIOR APPLICATION NUMBER: 60/427090
PRIOR APPLICATION NUMBER: 60/427090
PRIOR APPLICATION NUMBER: 60/427090
PRIOR FILING DATE: 2003-11-15
NUMBER OF SEQ ID NOS: 372
SEQ ID NO 343
LENGTH: 19
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US-10-714-195-344/C

Sequence 344, Application US/10714195

Publication No. US20050019785A1

GENERAL INFORMATION:

APPLICANT: Baker, Joffre

APPLICANT: Cronin, Maureen

APPLICANT: Cronin, Maureen

APPLICANT: Baselga, Jose

TITLE OF INVENTION: GENE EXPRESSION PROFILING OF EGFR

TITLE OF INVENTION: POSITIVE CANCER

FILE REFERENCE: 39740-0005
                                                                                                                                                                                                                                                                                                                                                                                   Query Match
4.2%; Score 19; DB
Best Local Similarity 100.0%; Pred. No. 82;
Matches 19; Conservative 0; Mismatches
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4.2%; Score 19; DB
Best Local Similarity 100.0%; Pred. No. 82;
Matches 19; Conservative 0; Mismatches
                PRIOR APPLICATION NUMBER: US 60/345,326
PRIOR FILING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: US 60/359,196
PRIOR APPLICATION NUMBER: US 60/383,195
PRIOR APPLICATION NUMBER: US 60/383,195
PRIOR FILING DATE: 2002-05-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: PatentIn version 3.2
SEQ ID NO 19
LENGTH: 19
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PRIOR FILING DATE: 2002-10-16
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                                                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Artificial
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; OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/siNA sense re
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4.2%; Score 19; DB 1; Length 19;
Best Local Similarity 100.0%; Pred. No. 82;
Matches 19; Conservative 0; Mismatches 0; Indels
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4.2%; Score 19; DB 1; Length 19;
Best Local Similarity 78.9%; Pred. No. 82;
Matches 15; Conservative 4; Mismatches 0; Indels
CURRENT APPLICATION NUMBER: US/10/714,195
CURRENT FILING DATE: 2003-11-14
PRIOR APPLICATION NUMBER: 60/427090
PRIOR FILING DATE: 2003-11-15
NUMBER OF SEQ ID NOS: 372
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 344
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              431 CAGGACTCGGCTCACACAT 449
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                                                                                                                                                                                                                                                                                                                                            TYPE: DNA ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/#1NA sense re
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APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: Bagelahan, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400/209 (MRHBEN2-708-C)
CURRENT FILING DATE: 2004-08-20
CURRENT FILING DATE: 2004-08-20
FRIOR PAPLICATION NUMBER: US 60/396,600
FRIOR APPLICATION NUMBER: US 60/396,600
FRIOR APPLICATION NUMBER: US 10/826,966
FRIOR PELING DATE: 2004-01-14
FRIOR PELING DATE: 2004-01-14
FRIOR PELING DATE: 2004-01-14
FRIOR PELING DATE: 2003-11-24
FRIOR APPLICATION NUMBER: US 10/757,803
FRIOR PELING DATE: 2003-11-24
FRIOR PELING DATE: 2003-01-01-14
FRIOR PELING DATE: 2003-01-02-16
FRIOR APPLICATION NUMBER: US 10/444,853
FRIOR APPLICATION NUMBER: US 10/444,853
FRIOR APPLICATION NUMBER: US 10/444,853
FRIOR APPLICATION NUMBER: PET/US03/05346
FRIOR APPLICATION NUMBER: PET/US03/05028
FRIOR APPLICATION NUMBER: PET/US03-02-20
FRIOR APPLICATION NUMBER: PET/US03-02-20
FRIOR APPLICATION NUMBER: PET/US03-02-20
FRIOR APPLICATION NUMBER: PET/US03-02-20
FRIOR FILING DATE: 2003-02-20
FRIOR APPLICATION NUMBER: PET/US03/05028
FRIOR FILING DATE: 2003-02-20
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                                                                                   PRIOR APPLICATION NUMBER: US 10/85,966
PRIOR FILING DATE: 2004-04-16
PRIOR PILING DATE: 2004-04-16
PRIOR PILING DATE: 2004-04-16
PRIOR PILING DATE: 2004-04-16
PRIOR FILING DATE: 2004-01-124
PRIOR PELING DATE: 2003-11-24
PRIOR APPLICATION NUMBER: US 10/44,853
PRIOR FILING DATE: 2003-11-23
PRIOR PELING DATE: 2003-05-23
PRIOR APPLICATION NUMBER: PCT/US03/05346
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR FILING DATE: 2003-02-20
PRIOR PRIOR PRIOR PRIOR NUMBER: PCT/US03/05028
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PCT/US04/16390
                                                       FILING DATE: 2004-05-24
APPLICATION NUMBER: US 10/826,966
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             50 CCCTAACTGAGAAGGGCGT 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin version 3.3 SEQ ID NO 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity 84.2
Matches 16; Conservative
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US-10-923-330-8
                                                                                                                                                                                                                              US-10-922-330-8

US-10-922-330-8

Sequence 8, Application US/10923330

PUDIcation No. US20050153916A1

GENERAL INFORMATION:

APPLICANT: SITRA Therapeutics, Inc.

APPLICANT: McSWAGGEN, James

APPLICANTON NUMBER: US/10/923,330

CURRENT APPLICATION NUMBER: US 60/396,600

FRIOR FILING DATE: 2002-07-17

FRIOR FILING DATE: 2002-07-17

FRIOR FILING DATE: 2002-07-14

FRIOR FILING DATE: 2003-01-14

FRIOR FILING DATE: 2003-11-23

FRIOR FILING DATE: 2003-05-23

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; Sequence 9, Application US/10923330
; Publication No US20050153916A1
; GENERAL INFORMATION:
   APPLICANT: Strna Therapeutics, Inc.
   APPLICANT: McSwiggen, James
; APPLICANT: Beigelman, Leonid
; TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
; TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
; TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
; TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
; CURRENT APPLICATION NUMBER: US/10/923,330
; CURRENT FILING DATE: 2004-08-20
; PRIOR FILING DATE: 2004-05-24
; PRIOR FILING DATE: 2004-05-24
; PRIOR FILING DATE: 2002-07-17
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     1 UGGGCCUGGGAGGGGU 19
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               -10-923-330-9
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; OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/sinA sense re
US-10-923-330-12
                                                                                                       APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: BAY Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT PELICATION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-06-20
FRIOR PELICATION NUMBER: PCT/US03/04088
PRIOR FILING DATE: 2004-05-24
FRIOR PELICATION NUMBER: US 60/396,600
FRIOR PELICATION NUMBER: US 60/396
FRIOR PELICATION NUMBER: US 10/757,803
FRIOR PELICATION NUMBER: US 10/750,448
FRIOR PELICATION NUMBER: US 10/720,448
FRIOR PELICATION NUMBER: US 10/444,853
FRIOR PELICATION NUMBER: US 10/444,853
FRIOR PELICATION NUMBER: US 10/730,3059
FRIOR PELICATION NUMBER: US 10/730,3059
FRIOR PELICATION NUMBER: US 2003-01-23
FRIOR PELICATION NUMBER: US 2003-01-23
FRIOR PELICATION NUMBER: US 10/730,3050
FRIOR PELICATION NUMBER: US 2003-05-20
FRIOR PELICATION NUMBER: US 10/303-05-30
FRIOR P
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Sequence 13, Application US/10923330

Sequence 13, Application US/10923330

PUBLIcation No. US20050153916A1

GENERAL INFORMATION:

APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: Beigadman, Leonid

TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

CURRENT FILING DATE: 2004-08-20

FRIOR PELICATION NUMBER: US 10/923,330

CURRENT FILING DATE: 2004-06-24

PRIOR PLICATION NUMBER: US 60/396,600

PRIOR APPLICATION NUMBER: US 10/826,966

PRIOR FILING DATE: 2004-06-24

PRIOR FILING DATE: 2004-04-16

PRIOR PLING DATE: 2004-04-16

PRIOR PLING DATE: 2004-04-16

PRIOR PLING DATE: 2004-01-14

PRIOR PLING DATE: 2004-01-14
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4.2%; Score 19; DB 1; Length 19;
Best Local Similarity 73.7%; Pred. No. 82;
Matches 14; Conservative 5; Mismatches 0; Indels
                    APPLICANT: Sirna Therapeutics, Inc. APPLICANT: McSwiggen, James APPLICANT: Beigelman, Leonid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       104 CTCGCTGACTTTCAGCGGG 122
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ORGANISM: Artificial Sequence
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                                                                                                           ; OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/siNA sense US-10-923-330-10
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US-10-923-330-11
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1 Sequence 11, Application US/10923330

1 Sequence 11, Application US/10923330

2 Publication No. US20050153916A1

3 PELICANT: Sirna Therapeutics, Inc.

APPLICANT: McSwiggen, James

APPLICANT: Beigelman, Leonid

1 TILE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

1 TILE OF INVENTION: Expression Using Short Interfering RNA (siNA)

1 TILE OF INVENTION: APPLICANT: A00/209 (MBHB02-708-C)
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CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-08-20
FRIOR PELICATION NUMBER: US/10/923,330
FRIOR PELICATION NUMBER: US/10/928
FRIOR PELING DATE: 2004-05-24
FRIOR PELING DATE: 2004-05-24
FRIOR PELING DATE: 2000-07-17
FRIOR PELING DATE: 2000-07-17
FRIOR APPLICATION NUMBER: PCT/US04/16390
FRIOR FILING DATE: 2004-04-16
FRIOR PELING DATE: 2004-04-16
FRIOR PELING DATE: 2004-01-12
FRIOR PELING DATE: 2004-01-12
FRIOR PELING DATE: 2003-11-24
FRIOR PILING DATE: 2003-11-24
FRIOR PELING DATE: 2003-11-23
FRIOR PELING DATE: 2003-11-23
FRIOR PELING DATE: 2003-05-23
FRIOR PELING DATE: 2003-05-23
FRIOR PILING DATE: 2003-05-20
FRIOR PILING DATE: 2003-05-20
FRIOR FILING DATE: 2003-05-30
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US-10-923-330-12
; Sequence 12, Application US/10923330
; Publication No. US20050153916A1
                                                                                                                                                                                                                                                                                                                                                                                                                                     68 TAGGCGCCGTGCTTTTGCT 86
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TYPE: RNA
ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 63.2
Matches 12; Conservative
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                                                                              FEATURE:
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Target Sequence/siNA sense r
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Publication No. US20050153916A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Baigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Rxpression Using Short Interfering RNA (sINA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
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84.2%; Pred. No. 82;
tive 3; Mismatches 0; Indels
       Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: Description of Artificial Sequence:
Score 19; DB 1;
Pred. No. 82;
5; Mismatches
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Sequence 15, Application US/10923330
Publication No. US20050153916A1
GENERAL INFORMATION:
                                                                                                                                                        140 GCCGCCTTCCACCGTTCAT 158
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                                                                                                                                                                                              Query Match
Best Local Similarity 73.7%;
Matches 14; Conservative
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ORGANISM: Artificial Seguence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 84.2*
Matches 16; Conservative
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US-10-923-330-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/siNA sense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 14, Application US/10921330

PUBLIcation No. US2050153916A1

GENERAL INFORMATION:

APPLICANT: Sirna Therapeutice, Inc.

APPLICANT: McSW-2094, James

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

FILE REFERENCE: 400/209 (MBHBG2-708-C)

CURRENT PELLING DATE: 2004-08-24

FRIOR FILING DATE: 2004-08-24

FRIOR PILING DATE: 2004-08-24

FRIOR PLILING DATE: 2004-05-24

FRIOR PLILING DATE: 2004-05-24

FRIOR PLILING DATE: 2004-04-16

FRIOR PLILING DATE: 2004-04-16

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FRIOR FILING DATE: 2004-01-14

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FRIOR FILING DATE: 2003-01-14

FRIOR FILING DATE: 2003-11-23

FRIOR FILING DATE: 2003-11-24

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FRIOR FILING DATE: 2003-01-14

FRIOR FILING DATE: 2003-11-24

FRIOR FILING DATE: 2003-01-23

FRIOR FILING DATE: 2003-11-24

FRIOR FILING DATE: 2003-11-24

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FRIOR FILING DATE: 2003-02-20

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PRIOR FILING DATE: 2003-11-24
PRIOR PELICATION NUMBER: US 10/693,059
PRIOR FILING DATE: 2003-11-23
PRIOR PELING DATE: 2003-11-23
PRIOR PELING DATE: 2003-05-23
PRIOR PELICATION NUMBER: PCT/US03/05346
PRIOR PELING DATE: 2003-02-20
PRIOR PILING DATE: 2003-02-30
PRIOR PILING DATE
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ORGANISM: Artificial Sequence
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SEQ ID NO 14
LENGTH: 19
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Matches 17; Conservative
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OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/siNA sense re
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US-10-923-330-18
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| Sequence 18, Application US/10923330
| Publication No. US20050153916A1
| GRNERAL INFORMATION: US20050153916A1
| APPLICANT: Sirna Therapeutics, Inc.
| APPLICANT: McSwigen, James
| APPLICANT: McSwigen, James
| APPLICANT: Beigelman, Leonid
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
| FILE REPRENCE: 400/209 (WHHED-108-C)
| CURRENT PILING DATE: 2004-08-20
| FRIOR PELICATION NUMBER: US/10/923,330
| CURRENT PILING DATE: 2004-08-20
| FRIOR PELICATION NUMBER: US 6/396,600
| PRIOR FILING DATE: 2002-07-17
| PRIOR FILING DATE: 2002-07-17
| PRIOR FILING DATE: 2004-04-16
| PRIOR FILING DATE: 2003-04-16
| PRIOR FILING DATE: 2003-05-20
| PRIOR FILING DATE: 2003-05-30
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                      PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR FILIMO DATE: 2003-02-20
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 766
SOFTWARE: Patentin version 3.3
SEQ ID NO 17.0
LENGTH: 19
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4.2%; Score 19; DB 1; Length 19;
Best Local Similarity 84.2%; Pred. No. 82;
Matches 16; Conservative 3; Mismatches 0; Indels
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Pred. No. 82;
3; Mismatches 0; Indels
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ORGANISM: Artificial Sequence
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Best Local Similarity 84.2%;
Matches 16; Conservative
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Publication No. US20050153916A1

GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James

TITLE OF INVENTION: RAY Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: Expression Using Short Interfering RNA (SiNA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (SiNA)

FILE REFERENCE: 400/209 (MBHB02-708-C)

CURRENT FILING DATE: 2004-08-20

PRIOR PLILING DATE: 2004-05-24

PRIOR PILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-04-16

PRIOR FILING DATE: 2004-04-16

PRIOR FILING DATE: 2004-04-16

PRIOR FILING DATE: 2004-04-16

PRIOR FILING DATE: 2003-11-24

PRIOR FILING DATE: 2003-11-24

PRIOR FILING DATE: 2003-11-23

PRIOR APPLICATION NUMBER: US 10/720,448

PRIOR FILING DATE: 2003-11-23

PRIOR APPLICATION NUMBER: US 10/720,448

PRIOR PILING DATE: 2003-11-23

PRIOR PILING DATE: 2003-11-23
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                               CURRENT FILING DATE: 2004-08-20

PRIOR PELING DATE: 2004-08-20

PRIOR PELING DATE: 2004-08-20

PRIOR PILING DATE: 2004-05-24

PRIOR APPLICATION NUMBER: PCT/US04/16390

PRIOR APPLICATION NUMBER: DCT/US04/16390

PRIOR APPLICATION NUMBER: US 10/826,966

PRIOR PILING DATE: 2004-04-16

PRIOR PILING DATE: 2004-04-16

PRIOR PILING DATE: 2004-04-16

PRIOR PILING DATE: 2004-01-14

PRIOR PILING DATE: 2003-11-23

PRIOR APPLICATION NUMBER: US 10/757,803

PRIOR PILING DATE: 2003-11-23

PRIOR PILING DATE: 2003-11-23

PRIOR PILING DATE: 2003-11-23

PRIOR PILING DATE: 2003-12-23

PRIOR PILING DATE: 2003-02-20

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PRIOR PILING DATE: 2003-02-20

PRIOR PILING DATE: 2003-02-20
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CURRENT APPLICATION NUMBER: US/10/923,330
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APPLICATION NUMBER: PCT/US03/05346
FILING DATE: 2003-02-20
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1 AUGUCAGCUGCUGGCCCGU 19
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ORGANISM: Artificial Sequence
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Matches 14; Conserv
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OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/#1NA gense re
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APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: McSwiggen, James

APPLICANT: McSwiggen, James

APPLICANT: McSwiggen, Jemes

APPLICANT: McSwiggen, Jemes

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Ge

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Ge

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Ge

TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

FILE PREPRENCE: 400/209 (MBHB02-708-C)

CURRENT FILING DATE: 2004-08-20

FRIOR PELING DATE: 2004-05-24

FRIOR PELING DATE: 2004-01-14

FRIOR PELING DATE: 2004-01-14

FRIOR PELING DATE: 2004-01-14

FRIOR PELING DATE: 2004-01-14

FRIOR PELING DATE: 2003-11-23

FRIOR PELING DATE: 2003-02-20

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                                               PRIOR FILING DATE: 2004-01-16
PRIOR PILING DATE: 2004-01-14
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PRIOR PILING DATE: 2003-11-24
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PRIOR PILING DATE: 2003-02-20
PRIOR PILING DATE: 2003-02-30
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                                           US 10/826,966
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SEQ ID NO 21
LENGTH: 19
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2004-05-24
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US-10-923-330-19
                                                                                                                                                         Sequence 19, Application US/10923330
| Publication No. US20050153916A1
| GENERAL INFORMATION: | GENERAL INFORMATION: | Publication No. US20050153916A1 | |
| APPLICANT: Sirma Therapeutics, Inc. | APPLICANT: MCSWIGGEN, Leonid | APPLICANT: MCSWIGGEN, Leonid | TITLE OF INVENTION: Expension Using Short Interfering RNA (sinA) |
| FILE REFERENCE: 400/209 (WBHBG2-708-C) | CURRENT APPLICATION NUMBER: US/10/923,330 |
| CURRENT APPLICATION NUMBER: US/10/923,330 |
| CURRENT APPLICATION NUMBER: US 00/3-6.00 |
| PRIOR APPLICATION NUMBER: US 00/3-6.00 |
| PRIOR PLING DATE: 2004-08-20 |
| PRIOR PLING DATE: 2004-04-16 |
| PRIOR PLING DATE: 2004-04-16 |
| PRIOR PLING DATE: 2004-04-16 |
| PRIOR PLING DATE: 2003-11-23 |
| PRIOR PLING DATE: 2003-11-24 |
| PRIOR PLING DATE: 2003-11-23 |
| PRIOR PLICATION NUMBER: US 10/53,059 |
| PRIOR PLICATION NUMBER: US 10/44,853 |
| PRIOR PLICATION NUMBER: US 10/44,853 |
| PRIOR PLICATION NUMBER: PCT/US03/05028 |
| PRIOR PLICATION NUMBER: PCT/US03/05028 |
| PRIOR PLING DATE: 2003-05-23 |
| PRIOR PLING DATE: 2003-05-20 |
| PRIOR PLING DATE: 2003-05-33 |
| PRIOR PLING DATE: 2003-05-30 |
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Sequence 20, Application US/20050153916A1

Publication No. US20050153916A1

GENERAL INFORMATION:

APPLICANT: Bisina Therapeutics, Inc.
APPLICANT: Beigelman, Leonid

TILLE OF INVENTION: RAN Interference Mediated Inhibition of Telomerase Gene
TILLE OF INVENTION: Expression Using Short Interfering RNA (BINA)

FILE REFERENCE: 400/209 (MBH802-708-C)

FURRENT APPLICATION NUMBER: US/310/203,330

CURRENT PELING DATE: 2004-06-20

PRIOR PILING DATE: 2004-05-24

PRIOR APPLICATION NUMBER: US 60/396,600

PRIOR FILING DATE: 2002-07-17

PRIOR PLING DATE: 2002-07-17

PRIOR APPLICATION NUMBER: PCT/US03/04/16390
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ORGANISM: Artificial Sequence
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LENGTH: 19
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Telomerase Gene

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OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/sinh sense re
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| Publication No US20050153916A1
| GENERAL INPORMATION:
| APPLICANT: Sirna Therapeutics, Inc.
| APPLICANT: Baigelman, Leconid
| TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
| TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
| FILE REPRENCE: 400/209 (MRH802-708-C)
| FILE REPRENCE: 2400/209 (MRH802-708-C)
| FILE REPRENCE: 2004-06-20
| FRIOR PLICATION NUMBER: DCT/US03/04088
| PRIOR FLILING DATE: 2004-05-24
| PRIOR FLILING DATE: 2004-06-24
| PRIOR FLILING DATE: 2004-06-24
| PRIOR FLILING DATE: 2004-06-24
| PRIOR FLILING DATE: 2004-06-14
| PRIOR FLILING DATE: 2004-01-14
| PRIOR FLILING DATE: 2003-11-24
                                                     APPLICANT: McSwiggen, James Hearles, Inc.
APPLICANT: McSwiggen, James Hearles, Inc.
APPLICANT: McSwiggen, James Hearles Hearle
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                            APPLICANT: Sirna Therapeutics, Inc.
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SEQ ID NO 23
LENGTH: 19
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 73.7
Matches 14; Conservative
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                                                                              FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/sinA sense
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Sequence 22, Application US/1092330

Publication No. US20050153916A1

GENERAL INFORMATION: The Messagen, James

APPLICANT: McSw4gen, James

FILE REFERENCE: 400/209 (MHHG2-708-C)

CURRENT FILING DATE: 2004-08-20

FRIOR APPLICATION NUMBER: US 6/396,600

FRIOR APPLICATION NUMBER: PCT/US03/0408

FRIOR FILING DATE: 2004-01-1

FRIOR FILING DATE: 2003-11-23

FRIOR FILING DATE: 2003-11-23

FRIOR FILING DATE: 2003-02-20

FRIOR FILING DATE: 2003-03-34

FRIOR FILING DATE: 2003-03-34
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4.2%; Score 19; DB
Best Local Similarity 84.2%; Pred. No. 82;
Matches 16; Conservative 3; Mismatches
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US-10-923-330-23
Sequence 23, Application US/10923330
Publication No. US20050153916A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                   266 CCGGGCTTCTCCGGAGGC 284
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TYPE: RNA
ORGANISM: Artificial Sequence
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Matches 18; Conserv
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US-10-923-330-21
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OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/sinA sense re
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APPLICANT: Birna Therapeutics, Inc.
APPLICANT: Birna Therapeutics, Inc.
APPLICANT: Begalama, James
APPLICANT: Begalama, Leonid
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
TITLE OF INVENTION: WHBER: US/10/923,330
CURRENT FILING DATE: 2004-08-20
FRIOR PELION OF Expression Using Space Short
FRIOR APPLICATION NUMBER: US 60/396,600
FRIOR APPLICATION NUMBER: US 60/396,600
FRIOR APPLICATION NUMBER: US 10/826,966
FRIOR FILING DATE: 2004-03-16
FRIOR FILING DATE: 2004-01-14
FRIOR PELION NUMBER: US 10/757,803
FRIOR APPLICATION NUMBER: US 10/757,803
FRIOR PELING DATE: 2003-11-24
FRIOR APPLICATION NUMBER: US 10/750,448
FRIOR FILING DATE: 2003-11-24
FRIOR FILING DATE: 2003-11-24
FRIOR FILING DATE: 2003-05-20
FRIOR FILING DATE: 2003-05-20
FRIOR FILING DATE: 2003-05-20
FRIOR APPLICATION NUMBER: US 10/703/05346
FRIOR FILING DATE: 2003-05-20
FRIOR APPLICATION NUMBER: PCT/US03/0508
FRIOR APPLICATION NUMBER: PCT/US03/0508
FRIOR FILING DATE: 2003-02-20
FRIOR APPLICATION NUMBER: PCT/US03/0508
FRIOR FILING DATE: 2003-02-20
FRIOR FILING DATE: 2003-02-20
FRIOR APPLICATION NUMBER: PCT/US03/0508
FRIOR FILING DATE: 2003-02-20
FRIOR APPLICATION NUMBER: PCT/US03/0508
FRIOR FILING DATE: 2003-02-20
FRIOR FILING DAT
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Publication No. US20050153916A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
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82;
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Best Local Similarity 84.2%; Pred. No. |
Matches 16; Conservative 3; Mismatch
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                                                                                                                                338 CGAGGGGGAGGTTCAGGCC 356
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                                                                                                                                                                             1 CGAGGGCGAGGUUCAGGCC 19
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                                                 17; Conservative
Best Local Similarity
Matches 17; Conserv
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US-10-923-330-27
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US-10-923-330-24
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US-10-923-330-25
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APPLICANT: Sirna Therapeutics, inc.
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
FILE REFERENCE: 400/209 (MRHBD2-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT APPLICATION NUMBER: PCT/US03/04088
PRIOR FILING DATE: 2004-08-20
PRIOR PILING DATE: 2002-07-17
PRIOR FILING DATE: 2002-07-17
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-01-4
PRIOR FILING DATE: 2004-01-14
PRIOR FILING DATE: 2004-01-14
PRIOR FILING DATE: 2003-11-24
PRIOR PLING DATE: 2003-11-24
PRIOR FILING DATE: 2003-11-24
PRIOR PLING DATE: 2003-11-24
PRIOR FILING DATE: 2003-02-20
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PRIOR APPLICATION NUMBER: US 10/693,059
PRIOR FILING DATE: 2003-11-23
PRIOR PILING DATE: 2003-11-23
PRIOR PLILING DATE: 2003-05-23
PRIOR PLILING DATE: 2003-05-23
PRIOR APPLICATION NUMBER: PCT/US03/05346
PRIOR PILING DATE: 2003-02-20
PRIOR PILING DATE: 2003-02-20
Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER: OF SEQ ID NOS: 768
SOFTWARE: Patentin version 3.3
SEQ ID NO 24
LENGTH: 19
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84.2%; Pred. No. 82;
tive 3; Mismatches 0; Indels
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; Sequence 25, Application US/10923330
; Publication No. US20050153916A1
; GENERAL INFORMATION:
; APPLICANT: Sirna Therapeutice, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  320 CCGCGGGTCTCTCGGGGGC 338
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: RNA
ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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Best Local Similarity
Matches 16; Conserva
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LENGTH: 19
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Score 19; DB 1; Length 19;

4.2%;

Query Match

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; OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/siNA sense re
US-10-923-330-28
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-08-20
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PRIOR APPLICATION NUMBER: DCT/US03/04088

PRIOR FILING DATE: 2004-05-24

PRIOR PELING DATE: 2002-07-17

PRIOR FILING DATE: 2002-07-17

PRIOR FILING DATE: 2004-05-24

PRIOR PLING DATE: 2004-05-24

PRIOR PLING DATE: 2004-05-07-07

PRIOR PLING DATE: 2004-05-07-07

PRIOR PLING DATE: 2004-05-07-07

PRIOR PLING DATE: 2004-05-07-07

PRIOR PLING DATE: 2004-01-14

PRIOR PLING DATE: 2004-01-14

PRIOR PLING DATE: 2003-11-23

PRIOR PLING DATE: 2003-11-23

PRIOR PLING DATE: 2003-05-23

PRIOR PLING DATE: 2003-05-23

PRIOR PLING DATE: 2003-05-23

PRIOR PLING DATE: 2003-02-20

PRIOR APPLICATION NUMBER: PCT/US03/05028

PRIOR PLING DATE: 2003-02-20

PRIOR PLING DATE: 2003-
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                                                                           Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER OF SEQ ID NOS: 768
SOFTWARE: PatentIn version 3.3
SEQ ID NO 28
LENGTH: 19
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Pred. No. 82;
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4.2%; Score 19; DB 1;
Best Local Similarity 78.9%; Pred. No. 82;
Matches 15; Conservative 4; Mismatches
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                                                                                                                                                                                                                                                                TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 89.5%;
Matches 17; Conservative
                                       PRIOR FILING DATE: 2003-02-20
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ORGANISM: Artificial Sequence
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Publication No. US20050153916A1

GENERAL INPORMATION:

APPLICANT: McSwiggen, James

APPLICANT: McSwiggen, James

APPLICANT: Beigelman, Leonid

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

FILE REFERENCE: 400/209 (MBHD2-708-C)

CURRENT APPLICATION NUMBER: US/10/23, 330

CURRENT APPLICATION NUMBER: US/10/23, 330

CURRENT APPLICATION NUMBER: US 60/396,600

PRIOR APPLICATION NUMBER: US 60/396,600

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24

PRIOR APPLICATION NUMBER: US 60/396,600

PRIOR APPLICATION NUMBER: US 10/85,966

PRIOR APPLICATION NUMBER: US 10/85,966

PRIOR APPLICATION NUMBER: US 10/720,448

PRIOR APPLICATION NUMBER: US 10/720,448

PRIOR PLING DATE: 2004-01-16

PRIOR APPLICATION NUMBER: US 10/720,448

PRIOR PLING DATE: 2003-11-24

PRIOR PLING DATE: 2003-02-20

PRIOR PLING DATE: 2003-02-20

PRIOR PLING DATE: 2003-02-20

PRIOR PRIOR PLING DATE: 2003-02-20
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NUMBER OF SEQ ID NOS: 768
SOFTWARE: Patentin version 3.3
SEQ ID NO 27
LENGTH: 19
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4.2%; Score 19; DB 1; Length 19;
Best Local Similarity 94.7%; Pred. No. 82;
Matches 18; Conservative 1; Mismatches 0; Indels
CURRENT FILING DATE: 2004-08-20
PRIOR APPLICATION NUMBER: PCT/US03/04088
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2002-07-17
PRIOR APPLICATION NUMBER: PCT/US04/16390
PRIOR FILING DATE: 2004-05-65
PRIOR FILING DATE: 2004-01-6
PRIOR PLILNG DATE: 2004-01-6
PRIOR PLILNG DATE: 2004-01-14
PRIOR PLILNG DATE: 2004-01-14
PRIOR PLILNG DATE: 2003-11-23
PRIOR APPLICATION NUMBER: US 10/757, 803
PRIOR PLILNG DATE: 2003-11-23
PRIOR PLILNG DATE: 2003-05-23
PRIOR PLING DATE: 2003-05-23
PRIOR PLING DATE: 2003-05-23
PRIOR PLING DATE: 2003-05-23
PRIOR PLILNG DATE: 2003-05-20
PRIOR PLILNG DATE: 2003-02-20
PRIOR PLILNG DATE: 2003-02-20
PRIOR PLILNG DATE: 2003-02-20
PRIOR PLILNG DATE: 2003-02-20
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; OTHER INFORMATION: Description of Artificial Sequence: Target Sequence/sinA sense re
US-10-923-330-31
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US-10-922-330-380.

US-10-922-330-380.

Publication No. US20050153916A1

GENERAL INFORMATION:

HORDWATTON:

APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: McSwiggen, James

MRIOR FILING DATE: 2004-08-20

FRIOR FILING DATE: 2004-01-14

FRIOR FILING DATE: 2003-11-23

FRIOR FILING DATE: 2003-05-22

FRIOR FILING DATE: 2003-05-23

FRIOR FILING DATE: 2003-05-3

FRIOR FILING DATE: 2003
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PRIOR FILLING DATE: 2004-01-16
PRIOR FILLING DATE: 2004-01-14
PRIOR PELING DATE: 2004-01-14
PRIOR PLING DATE: 2004-01-14
PRIOR FILING DATE: 2003-11-24
PRIOR FILING DATE: 2003-11-23
PRIOR PLICATION NUMBER: US 10/693,059
PRIOR FILING DATE: 2003-01-23
PRIOR FILING DATE: 2003-05-23
PRIOR FILING DATE: 2003-05-20
PRIOR FILING DATE: 2003-02-20
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84.2%; Pred. No. 82;
tive 3; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 84.2*
Matches 16; Conservative
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                                                                                                                                                                                                                                                 PAPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Microphyliolands
APPLICANT: Microphyliolands
APPLICANT: Microphyliolands
APPLICANT: Microphyliolands
APPLICANT: Microphyliolands
APPLICANT: Microphyliolands
TITLE OF INVENTION: Expression Underfering Short Interfering RNA (sinA)
FILE REPRENCE: 400/209 (MRH002-708-C)
FILE REPRENCE: 400/209 (MRH002-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT PELING DATE: 2004-08-20
FRIOR PRILING DATE: 2004-08-20
FRIOR PELICATION NUMBER: US 60/396,600
FRIOR PELICATION NUMBER: US 10/826,966
FRIOR FILING DATE: 2004-01-14
FRIOR FILING DATE: 2004-01-14
FRIOR FILING DATE: 2004-01-14
FRIOR FILING DATE: 2003-11-24
FRIOR FILING DATE: 2003-11-24
FRIOR FILING DATE: 2003-11-24
FRIOR FILING DATE: 2003-11-23
FRIOR FILING DATE: 2003-11-23
FRIOR FILING DATE: 2003-11-23
FRIOR PELICATION NUMBER: US 10/693,059
FRIOR FILING DATE: 2003-11-23
FRIOR FILING DATE: 2003-11-23
FRIOR APPLICATION NUMBER: PCT/US03/05286
FRIOR APPLICATION NUMBER: PCT/US03/05028
FRIOR APPLICATION NUMBER: FCT/US03/05028
FRIOR APPLICATION NUMBER: FCT/US03/05028
FRIOR APPLICATION NUMBER: FCT/US03/05028
FRIOR APPLICATION NUMBER: PCT/US03/05028
FRIOR APPLICATION NUMBER: PCT/US03/
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| Publication No. US20050153916A1
| Publication No. US20050153916A1
| GENERAL INFORMATION:
| APPLICANT: Strna Therapeutics, Inc.
| APPLICANT: Beiggelman, Leonid
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
| TITLE OF INVENTION NUMBER: US/10/923,330
| PRIOR FILING DATE: 2004-06-24
| PRIOR FILING DATE: 2004-06-24
| PRIOR FILING DATE: 2004-05-24
| PRIOR FILING DATE: 2004-05-24
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                                                                                                                               Sequence 30, Application US/10923330 Publication No. US20050153916A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       428 ACCCAGGACTCGGCTCACA 446
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ORGANISM: Artificial Sequence
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US-10-923-330-31
                                                                                         US-10-923-330-30
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| Publication No. US20050153916A1
| Sequence 39, Application US/1092330
| Publication No. US20050153916A1
| Sequence 39, Application US20050153916A1
| SEQUENCE STITE OF UNIVERTION: MCMSQGEN, James
| APPLICANT: MCSWAGOEN, James
| PAPLICANT: MCSWAGOEN, James
| APPLICANT: MCSWAGOEN, James
| PAPLICANT: MCSWAGOEN, James
| PAPLICANT: MCSWAGOEN, JAMES US 00/209 (MRH02-700-C)
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sink)
| FILE REFERENCE: 400/209 (MRH02-700-C)
| FILE PEPLICATION NUMBER: US/10/923,330
| PRIOR APPLICATION NUMBER: US 00/396,600
| PRIOR PELING DATE: 2004-05-24
| PRIOR PELING DATE: 2004-05-34
| PRIOR FLING DATE: 2004-01-14
| PRIOR FLING DATE: 2004-01-14
| PRIOR PLING DATE: 2004-01-14
| PRIOR PLING DATE: 2003-11-23
| PRIOR PLING DATE: 2003-11-23
| PRIOR FLING DATE: 2003-05-23
| PRIOR PLING DATE: 2003-05-24
| PRIOR PLING DATE:
                               FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
US-10-923-330-38
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4.2%; Score 19; DB 1; Length 19;
Best Local Similarity 100.0%; Pred. No. 82;
Matches 19; Conservative 0; Mismatches 0; Indels
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4.2%; Score 19; DB 1; Length 19;
Best Local Similarity 100.0%; Pred. No. 82;
Matches 19; Conservative 0; Mismatches 0; Indels
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US-10-923-330-40/c
Sequence 40, Application US/10923330;
Publication No. US20050153916A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
                                                                                                                                                                                                                                                                                                                                   14 TGGGCCTGGGAGGGGTGGT 32
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ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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US-10-923-330-39/c
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RESULT 108

US-10-23-330-42/c

SQUEUTIC 42, Application US/1022330

Fublication No. US20050153916A1

SEQUENCE 42, Application US/1022330

Fublication No. US20050153916A1

SAPLICANT: SATE Therapeutics, Inc.

APPLICANT: Beigelman, Leonid

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RNA INTERFERENCE: 400/209 (MRH802-708-C)

CURRENT APPLICATION NUMBER: US 400/209 (MRH802-0)

FRIOR FILING DATE: 2004-06-24

FRIOR FILING DATE: 2004-06-24

FRIOR APPLICATION NUMBER: US 60/396,600

FRIOR FILING DATE: 2004-01-07-17

FRIOR APPLICATION NUMBER: US 10/720,448

FRIOR APPLICATION NUMBER: US 10/720,448

FRIOR PLING DATE: 2004-01-124

FRIOR FILING DATE: 2004-01-124

FRIOR PLING DATE: 2003-11-24

FRIOR FILING DATE: 2003-11-24

FRIOR PLING DATE: 2003-11-24

FRIOR PLING DATE: 2003-11-24

FRIOR PLING DATE: 2003-11-23

FRIOR APPLICATION NUMBER: US 10/593,059

FRIOR PLING DATE: 2003-11-23

FRIOR PLING DATE: 2003-10-24

FRIOR PREPARENTE PREP
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PRIOR FILING DATE: 2003-11-23
PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR FILING DATE: 2003-05-23
PRIOR FILING DATE: 2003-02-20
PRIOR FILING DATE: 2003-02-20
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER: Patentin version 3.3
SEQ ID NOS: 768
SOFTWARE: Patentin version 3.3
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: RNA
ORGANISM: Artificial Sequence
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Best Local Similarity 100.0
Matches 19; Conservative
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4.2%; Score 19; DB 1; Length 19; 100.0%; Pred. No. 82;

Query Match Best Local Similarity

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GENERAL INFORMATION:
APPLICANT: BITAGE AND AND AND APPLICANT: BITAGE AND APPLICANT: BEADLICANT: BOOK AND AND ADDITION: RAY INTERFERENCE: 400/209 (MRHBDA2-708-C)
FILE OF INVENTION: EXPRESSION USING Short Interfering RNA (siNA)
FILE OF INVENTION: RAY BEEN COLOR - 100 (MRHBDA2-708-C)
CURRENT FILING DATE: 2004-08-20
FRIOR PAPLICATION NUMBER: US 10/396,600
FRIOR PAPLICATION NUMBER: US 60/396,600
FRIOR APPLICATION NUMBER: US 10/396,600
FRIOR PELING DATE: 2004-05-24
FRIOR PELING DATE: 2004-05-24
FRIOR PELING DATE: 2004-01-16
FRIOR PELING DATE: 2004-01-16
FRIOR PELING DATE: 2004-01-14
FRIOR PELING DATE: 2003-11-23
FRIOR PELING DATE: 2003-11-24
FRIOR PELING DATE: 2003-11-24
FRIOR PELING DATE: 2003-05-20
FRIOR FILING DATE: 2003-05-20
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Publication No. US20050153916A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Beigelman, Leonid
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-08-20
Gaps
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Indels
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0; Mismatches
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                                                                          86 TCCCCGCGCGCTGTTTTC 104
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                                                                                                                 19 TCCCCGCGCGCTGTTTTC 1
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Conservative
                                                                                                                                                                                                                                                                                            -10-923-330-43/c
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19;
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Page 44

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PUDLICATION NO. USZOUSO153916A1

APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-08-20
FRIOR APPLICATION NUMBER: US 60/396,600
FRIOR PLING DATE: 2004-06-24
FRIOR PLING DATE: 2004-06-24
FRIOR PLING DATE: 2004-06-24
FRIOR PLING DATE: 2004-01-14
FRIOR APPLICATION NUMBER: US 10/693,059
FRIOR APPLICATION NUMBER: US 10/693,059
FRIOR APPLICATION NUMBER: US 10/693,059
FRIOR PLING DATE: 2003-11-24
FRIOR APPLICATION NUMBER: US 10/693,059
FRIOR PLING DATE: 2003-01-13
FRIOR PLING DATE: 2003-02-20
FRIOR PLING DATE: 2003-02-20
FRIOR APPLICATION NUMBER: US 10/693,059
FRIOR PLING DATE: 2003-02-20
FRIOR PLING DATE: 2003-02-20
FRIOR APPLICATION NUMBER: US 10/693,050
FRIOR PLING DATE: 2003-02-20
FRIOR PLING DATE: 2003-02-20
FRIOR FILING DATE: 2003-03-20-20
FRIOR FILING DATE: 2003-03-2
                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region US-10-923-330-45
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Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER OF SEQ ID NOS: 768 SOFTWARE: PatentIn version 3.3 SEQ ID NO 45
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Best Local Similarity 100.0%; Pred. No. 82;
Matches 19; Conservative 0; Mismatches
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ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
                                                                                                                                                                                    LENGTH: 19
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1 US-10-923-3310-45/C

1 US-10-923-3310-45/C

2 Sequence 45, Application US/10923330

3 Publication No. US20050153916A1

3 GENERAL INFORMATION:

4 APPLICANT: Sirna Therapeutics, Inc.

5 APPLICANT: Beigelman, Leonie

5 TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

7 TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

7 TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

7 TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

7 TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

7 TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

7 TITLE OF INVENTION WHERE: US/10/923,330

7 CURRENT APPLICATION WHERE: US (0/396,600

7 FRIOR FILING DATE: 2004-05-24

7 FRIOR FILING DATE: 2004-05-24

7 FRIOR FILING DATE: 2004-05-24

7 FRIOR FILING DATE: 2004-01-14

7 FRIOR FILING DATE: 2003-11-24

7 FRIOR FILING DATE: 2003-11-24

7 FRIOR FILING DATE: 2003-11-23

7 FRIOR FILING DATE: 2003-11-23

7 FRIOR FILING DATE: 2003-01-123

7 FRIOR FILING DATE: 2003-01-123

7 FRIOR FILING DATE: 2003-02-20

7 FRIOR FILING DATE: 2003-02-20
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                                                       ## FRIOR APPLICATION NUMBER: US 60/396,600
#PRIOR FILING DATE: 2002-07-17
#PRIOR FILING DATE: 2002-07-17
#PRIOR FILING DATE: 2004-05-24
#PRIOR PELING DATE: 2004-05-05
#PRIOR PELING DATE: 2004-05-05
#PRIOR PELING DATE: 2004-04-16
#PRIOR APPLICATION NUMBER: US 10/757,803
#PRIOR PILING DATE: 2004-01-14
#PRIOR PILING DATE: 2003-11-24
#PRIOR PILING DATE: 2003-11-24
#PRIOR PILING DATE: 2003-11-23
#PRIOR APPLICATION NUMBER: US 10/444,853
#PRIOR APPLICATION NUMBER: PC1/US03/05946
#PRIOR APPLICATION NUMBER: PC1/US03/05346
#PRIOR PILING DATE: 2003-02-23
#PRIOR PILING DATE: 2003-02-20
#PRIOR PILING DATE: 2003-02-20
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Pred. No. 82;
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    APPLICATION NUMBER: PCT/US03/04088
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ORGANISM: Artificial Sequence
                                                   2004-05-24
                                          FILING DATE: 2004-09
APPLICATION NUMBER:
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Matches 19
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   GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400/209 (MBHB02-709/23),330
CURRENT APPLICATION NUMBER: PCT/US03/04088
PRIOR PILING DATE: 2004-05-24
PRIOR APPLICATION NUMBER: US 60/396,600
PRIOR APPLICATION NUMBER: PCT/US04/16390
PRIOR APPLICATION NUMBER: PCT/US04/16390
PRIOR APPLICATION NUMBER: US 10/826,966
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4.2%; Score 19; DB 1; Length 19;
Best Local Similarity 100.0%; Pred. No. 82;
Matches 19; Conservative 0; Mismatches 0; Indels
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ORGANISM: Artificial Sequence
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Beigelman, Jemes
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REPRESENCE: 400/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: BC1/0923,330
CURRENT FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-06-24-16
PRIOR FILING DATE: 2004-04-16
PRIOR FILING DATE: 2004-04-16
PRIOR FILING DATE: 2004-01-14
PRIOR FILING DATE: 2004-01-14
PRIOR FILING DATE: 2004-01-14
PRIOR FILING DATE: 2004-11-23
PRIOR FILING DATE: 2003-11-24
PRIOR FILING DATE: 2003-11-23
PRIOR FILING DATE: 2003-11-23
PRIOR FILING DATE: 2003-11-23
PRIOR FILING DATE: 2003-05-33
PRIOR PRIOR PRIOR OWNER: PCT/US03/05028
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR PRIOR PRIOR OFFICE 2003-02-20-20
PRIOR PRIOR PRIOR OFFICE 2003-02-20-20
PRIOR PRIOR PRIOR DATE: 2003-02-20
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                                  PRIOR PILING DATE: 2004-01-14
PRIOR PILING DATE: 2003-11-24
PRIOR PILING DATE: 2003-11-24
PRIOR PILING DATE: 2003-11-24
PRIOR PILING DATE: 2003-11-23
PRIOR PILING DATE: 2003-11-23
PRIOR PELING DATE: 2003-11-23
PRIOR PELING DATE: 2003-05-23
PRIOR PELING DATE: 2003-05-20
PRIOR PILING DATE: 2003-02-20
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4.2%; Score 19; DB
Best Local Similarity 100.0%; Pred. No. 82;
Matches 19; Conservative 0; Mismatches
PLICATION NUMBER: US 10/757,803
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ORGANISM: Artificial Sequence
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| Publication No. US20050153916A1
| GENERAL INFORMATION:
| APPLICANT: Siria Therapeutics, Inc.
| APPLICANT: Siria Therapeutics, Inc.
| APPLICANT: Beisglamm, Leonid
| TITLE OF INVENTION: RNA Interference Mediated inhibition of Telomerase Gene
| TITLE OF INVENTION: RNA Interference Mediated inhibition of Telomerase Gene
| TITLE OF INVENTION: RNA Interference Mediated inhibition of Telomerase Gene
| TITLE OF INVENTION: RNA Interference Mediated inhibition of Telomerase Gene
| TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
| CURRENT APPLICATION NUMBER: US/10/223,330
| CURRENT APPLICATION NUMBER: US 60/396,600
| PRIOR PAPLICATION NUMBER: US 60/396,600
| PRIOR PAPLICATION NUMBER: US 10/486,524
| PRIOR PAPLICATION NUMBER: US 10/486,966
| PRIOR PLING DATE: 2004-04-16
| PRIOR PLING DATE: 2004-04-16
| PRIOR PLING DATE: 2004-01-14
| PRIOR PLING DATE: 2003-11-24
| PRIOR PLING DATE: 2003-11-24
| PRIOR PLING DATE: 2003-11-23
| PRIOR PLING DATE: 2003-11-23
| PRIOR PLING DATE: 2003-05-23
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
US-10-923-330-49
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SOFTWARE: PatentIn version 3.3
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                                                                                                                                Query Match 4.2%; Score 19; DB Best Local Similarity 100.0%; Pred. No. 82; Matches 19; Conservative 0; Mismatches
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US-10-923-330-51/c
; Sequence 51, Application US/10923330
; Publication No. US2050153916A1
; GENERAL INFORMATION:
; APPLICANT: Sirna Therapeutics, Inc.
; APPLICANT: McSwiggen, James
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ORGANISM: Artificial Sequence
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Matches 19; Conserval
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                                                                                        -10-923-330-54/c
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US-10-923-330-53
                                                                                                                                                                             OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
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100.0%; Pred. No. 82;
tive 0; Mismatches 0; Indels
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Pred. No. 82;
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PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR FILING DATE: 2003-05-23
PRIOR APPLICATION NUMBER: PCT/US03/05346
PRIOR FILING DATE: 2003-02-20
PRIOR PLING DATE: 2003-02-20
PRIOR PLING DATE: 2003-02-20
Remaining Prior Application data removed - 8
SOFTWARE: Patentin version 3.3
SEQ ID NO 52
                                                                                                                                                                                                                                                                    266 CCGGGGCTTCTCCGGAGGC 284
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                                                                                                                                          TYPE: RNA
ORGANISM: Artificial Sequence
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Best Local Similarity 100.
Matches 19; Conservative
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GENERAL INCOMMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Beigelman, Leonid
ITILLE OF INVENTION RAN Interference Mediated Inhibition of Telomerase Gene
ITILLE OF INVENTION RAN Interference Mediated Inhibition of Telomerase Gene
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ITILE OF INVENTION RAN INCOMEN:
ITILE OF INVENTION WHERE: US/10/923,330
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT APPLICATION NUMBER: DOUG-08-20
PRIOR APPLICATION NUMBER: DOUG-24 16 900
PRIOR APPLICATION NUMBER: US 60/396,600
PRIOR PAPLING DATE: 2004-01-17
PRIOR PAPLICATION NUMBER: US 10/826,966
PRIOR PELING DATE: 2004-01-14
PRIOR PELING DATE: 2004-01-14
PRIOR PELING DATE: 2004-01-14
PRIOR PELING DATE: 2004-01-14
PRIOR PELING DATE: 2003-01-12
PRIOR PELING DATE: 2003-01-12
PRIOR PELING DATE: 2003-01-12
PRIOR PELING DATE: 2003-01-13
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PRIOR PELING DATE: 2003-01-13
PRIOR PELING DATE: 2003-02-20
PRIOR PELING DATE: 2003-03-20
PRIOR PELING DATE: 2003-03-30
PRIOR PELING DATE: 2003-
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Publication No. US20050153916A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: Rapression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-08-20
PRIOR APPLICATION NUMBER: PCT/US03/04088
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Sequence 54, Application US/10923330 Publication No. US20050153916A1 GENERAL INFORMATION:
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Gaps

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PULICARION NO. USZUOSOIS3916A1

| PULICARION NO. USZUOSOIS3916A1
| APPLICANT: Sirna Therapeutics, Inc.
| APPLICANT: Bajgalman, Lecond |
| TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Generation of TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Generation Procession Using Short Interfering RNA (siNA)
| TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA) |
| TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA) |
| TITLE OF INVENTION: UNPRER: US/10/923,330 |
| CURRENT FILING DATE: 2004-08-20 |
| PRIOR APPLICATION NUMBER: US 60/396,600 |
| PRIOR PELING DATE: 2002-07-14 |
| PRIOR PELING DATE: 2004-06-24 |
| PRIOR PELING DATE: 2004-06-24 |
| PRIOR PELING DATE: 2004-01-14 |
| PRIOR PELING DATE: 2004-01-14 |
| PRIOR PELING DATE: 2004-01-12 |
| PRIOR PELING DATE: 2004-01-12 |
| PRIOR PELING DATE: 2003-01-24 |
| PRIOR PELING DATE: 2003-02-20 |
| PRIOR PELING DATE: 2003-02-03 |
| PRIOR PELING DATE: 2003-02-20 |
| PRIOR PELING DATE: 2003-02-03 |
| PRIOR PELING DATE: 2003-03-03 |
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Best Local Similarity 100.0%; Pred. No. 82;
Matches 19; Conservative 0; Mismatches
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                 NUMBER OF SEQ ID NOS: 768
SOFTWARE: Patentin version 3.3
SEQ ID NO 56
LENGTH: 19
                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence FEATURE:
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ORGANISM: Artificial Sequence
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Best Local Similarity
Matches 19; Conserva
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US-10-923-330-57/c
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APPLICANT: SITE A TREADELLES, INC.
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA INTERFERENCE: 400/209 (MBHB02-708-C)
CURRENT PELLING DATE: 2004-08-20
PRIOR PELLING DATE: 2004-05-24
PRIOR APPLICATION NUMBER: PCT/US04/16390
PRIOR FILING DATE: 2004-06-24
PRIOR PELLING DATE: 2004-06-24
PRIOR PELLING DATE: 2004-06-24
PRIOR PELLING DATE: 2004-06-24
PRIOR PELLING DATE: 2004-06-24
PRIOR FILING DATE: 2004-01-14
PRIOR PELLING DATE: 2004-01-14
PRIOR PELLING DATE: 2004-01-14
PRIOR PELLING DATE: 2003-11-23
PRIOR PELLING DATE: 2003-11-24
PRIOR PELLING DATE: 2003-11-23
PRIOR PELLING DATE: 2003-05-23
PRIOR PELLING DATE: 2003-05-23
PRIOR PELLING DATE: 2003-02-20
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PRIOR FILING DATE: 2004-05-24

PRIOR APPLICATION NUMBER: US 60/396,600

PRIOR APPLICATION NUMBER: US 60/396,600

PRIOR PILING DATE: 2002-07-17

PRIOR PILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-04-16

PRIOR PILING DATE: 2004-04-16

PRIOR PELING DATE: 2004-04-16

PRIOR PELING DATE: 2004-01-14

PRIOR APPLICATION NUMBER: US 10/720,448

PRIOR PILING DATE: 2003-11-24

PRIOR PILING DATE: 2003-11-23

PRIOR PILING DATE: 2003-11-23

PRIOR PILING DATE: 2003-12-3

PRIOR PILING DATE: 2003-05-23

PRIOR PILING DATE: 2003-02-20

PRIOR APPLICATION NUMBER: PCT/US03/05028

PROPER PILING DATE: 2003-02-20

PRIOR APPLICATION NUMBER: PCT/US03/05028

PROOR THING DATE: 2003-02-20

PRIOR APPLICATION NUMBER: PCT/US03/05028

PROPER PILING DATE: 2003-02-20

PRIOR PILING DATE: 2003-02-20

PRIOR APPLICATION NUMBER: PCT/US03/05028

PROPER PILING DATE: 2003-02-20

PRIOR PILING DATE: 2003-02-20
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APPLICANT: Sirna Therapeutics, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           320 CCGCGGGTCTCTCGGGGGC 338
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ORGANISM: Artificial Sequence
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; OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region US-10-923-330-57 ö Gaps . 0

RESULT 124

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; OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region US-10-922-330-59
PRIOR APPLICATION NUMBER: US 10/757,803

PRIOR FILING DATE: 2004-01-14

PRIOR APPLICATION NUMBER: US 10/720,448

PRIOR FILING DATE: 2003-11-24

PRIOR FILING DATE: 2003-11-23

PRIOR FILING DATE: 003-31-23

PRIOR FILING DATE: 2003-10-23

PRIOR FILING DATE: 2003-05-23

PRIOR PRIOR APPLICATION NUMBER: PCT/US03/05346

PRIOR APPLICATION NUMBER: PCT/US03/05028

PRIOR FILING DATE: 2003-02-20

PRIOR PLING DATE: 2003-02-20

Remaining Prior Application data removed - See File Wrapper or PALM.

SEQ ID NOS: 768

SEQ ID NOS: 768

LENGTH 19

LENGTH: 19

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ORGANISM: Artificial Sequence
FEATURE:
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US-10-923-330-59/c

i Sequence 59, Application US/10923330

publication No. US20050153916A1

GENERAL INFORMATION:

APPLICANT: Sirna Therapeutica, Inc.

APPLICANT: Beigalman, Leonid

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

FILE REFERENCE: 400/209 (MBHB02-708-C)

CURRENT FILING DATE: 2004-08-20

PRIOR PLICATION NUMBER: DC7/US03/04088

PRIOR FILING DATE: 2004-08-20

PRIOR PLICATION NUMBER: US 60/396,600

PRIOR APPLICATION NUMBER: US 60/396,600

PRIOR PLICATION NUMBER: US 60/396,600

PRIOR PLILON DATE: 2004-05-24

PRIOR PLING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-06-26

PRIOR FILING DATE: 2004-06-16
                                                                                                                                                             APPLICANT: SITRA interpention, int.

APPLICANT: Baigelman, Leonid

APPLICANT: Beigelman, Leonid

TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

FILE REPRENCE: 400/209 (MBHB02-708-C)

CURRENT APPLICATION NUMBER: US/10/923,330

CURRENT APPLICATION NUMBER: US/10/923,330

PRIOR APPLICATION NUMBER: US 60/396,600

PRIOR PLING DATE: 2004-06-20

PRIOR PLING DATE: 2004-06-21

PRIOR PLING DATE: 2004-06-24

PRIOR PLING DATE: 2004-06-24

PRIOR PLING DATE: 2004-01-14

PRIOR PLING DATE: 2004-01-14

PRIOR PLING DATE: 2004-01-14

PRIOR PLING DATE: 2004-01-14

PRIOR PLING DATE: 2003-01-1-24

PRIOR PLING DATE: 2003-01-24

PRIOR PLING DATE: 2003-01-23

PRIOR PLING DATE: 2003-01-23

PRIOR PLING DATE: 2003-02-20

PRIOR PLING DATE: 2003-02-3

PRIOR PLING DATE: 2003-02-3

PRIOR PLING DATE: 2003-02-3

PRIOR PLING DATE: 2003-02-3

PRIOR PLING DATE: 2003-02-20

PRIOR PLING DATE: 2003-02-20
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                                                                Sequence 58, Application US/10923330; Publication No. US20050153916A1; GENERAL INFORMATION: APPLICANT: Sirna Therapeutics, Inc.
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ORGANISM: Artificial Seguence
                                           US-10-923-330-58/c
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ORGANISM: Artificial Sequence
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Matches 19; Conservative
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UG-10-92-130-61/C

Sequence 61, Application US/10923330

Publication No. US2005013916A1

SEGNERAL INFORMATION:

APPLICANT: MCGRAPION:

APPLICANT: MCGRAPION:

APPLICANT: MCGRAPION:

APPLICANT: MCGRAPION:

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

FILE REPERENCE: 400/209 (HRHBOZ-708-C)

CURRENT PELICATION NUMBER: US/04-08-B

PRIOR PELICATION NUMBER: US 10/0923,330

CURRENT PELING DATE: 2004-06-24

PRIOR APPLICATION NUMBER: US 10/204/16390

PRIOR APPLICATION NUMBER: US 10/204/16390

PRIOR PELING DATE: 2004-01-14

PRIOR PELING DATE: 2004-01-14

PRIOR PELING DATE: 2003-11-24

PRIOR PELING DATE: 2003-11-24

PRIOR PELING DATE: 2003-11-24

PRIOR PELING DATE: 2003-11-24

PRIOR PELING DATE: 2003-02-20

PRIOR APPLICATION NUMBER: US 10/593,059

PRIOR APPLICATION NUMBER: US 10/593,059

PRIOR PELING DATE: 2003-02-20

PRIOR APPLICATION NUMBER: US 10/203/305346

PRIOR PELING DATE: 2003-02-20

PRIOR APPLICATION NUMBER: PCT/US03/05028

PRIOR FILING DATE: 2003-02-20

PRIOR PELING DATE: 2003-03-20-20

PRIOR PELING DATE: 2003-03-20-20

PRIOR PELING DATE: 2003-03-20-20

PRIOR PELING DATE: 2003-03-20-20

PRIOR PELING DATE: 200
OTHER INFORMATION: Description of Artificial Sequence: siNA antisense region
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82;
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4.2%; Score 19; DB 1; Length 19;
Best Local Similarity 100.0%; Pred. No. 82;
Matches 19; Conservative 0; Mismatches 0; Indels
                                                                                                  Query Match
4.2%; Score 19; DB
Best Local Similarity 100.0%; Pred. No. 82;
Matches 19; Conservative 0; Mismatches
                                                                                                                                                                                                                                            410 CTGAGCTGTGGGACGTGCA 428
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Publication No. US20050153916A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           428 ACCCAGGACTCGGCTCACA 446
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ORGANISM: Artificial Sequence
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              ; OTHER INFORM
US-10-923-330-60
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LENGTH: 19
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TITLE OF INVESTION: REAL INTEGRATION MARKED AND ADDRESS OF THE OFFICE OF INVESTION: REAL INTEGRATION PROPERTY OF THE OFFICE OF INVESTION: REAL INTEGRATION PROPERTY OF THE OFFICE OFFICE
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                         Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 100.0%; Pred. No. 1e+02; Matches 19; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                    Sequence 9, Application US/10831266
| Publication No. US2005003404A1
| GENERAL INFORMATION:
| APPLICANT: ROWLEY, Peter T.
| TITLE OF INVENTION: TELOMERASE INTERFERENCE TILE REFERENCE A-1266-1/RFT/THR
| CURRENT APPLICATION NUMBER: US/10/831,266
| CURRENT FILING DATE: 2004-04-22/RFILING DATE: 2001-10-16
| PRIOR APPLICATION NUMBER: US 60/345,326
| PRIOR FILING DATE: 2001-10-26
| PRIOR APPLICATION NUMBER: US 60/345,326
| PRIOR APPLICATION NUMBER: US 60/345,326
| PRIOR APPLICATION NUMBER: US 60/359,196
| PRIOR APPLICATION NUMBER: US 60/359,196
| PRIOR FILING DATE: 2002-02-20
| PRIOR APPLICATION NUMBER: US 60/383,195
| PRIOR PELING DATE: 2002-05-22
| NUMBER OF SEQ ID NOS: 17
| SOFTWARE: PALENTIN VERSION 3.2
| SEQ ID NO 9
| LENGTH: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 6, Application US/10831267
Fublication No. US2005009177A1
GENERAL INFORMATION:
APPLICANT: ROWLEY, Peter T.
TITLE OF INVENTION: TELOMERASE INTERFERENCE
FILE REFERENCE: A-71506-2/RFT/THR
CURRENT PELLOTION NUMBER: US/10/831,267
CURRENT FILING DATE: 2004-04-22
PRIOR PELLOTION NUMBER: US/0/33146
PRIOR PAPLICATION NUMBER: US 60/345,326
PRIOR APPLICATION NUMBER: US 60/345,326
PRIOR APPLICATION NUMBER: US 60/359,196
PRIOR APPLICATION NUMBER: US 60/359,196
PRIOR APPLICATION NUMBER: US 60/359,196
PRIOR PAPLICATION NUMBER: US 60/359,196
PRIOR APPLICATION NUMBER: US 60/339,195
PRIOR PELING DATE: 2002-05-22
NUMBER OF SEQ ID NOS: 23
SEQ ID NO 6
LENGTH: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: oligonucleotide US-10-831-266-9
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                                                                                                                                         270 GGCTTCTCCGGAGGCACCC 288
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ORGANISM: Artificial
FEATURE:
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Score 19; DB 1; Length 21;
Pred. No. 1e+02;
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4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches 0; Indels
                                                         0; Indels
                                                                                                                                                                                                                                                                Sequence 7, Application US/10831266

Sequence 7, Application US/10831266

Publication No. US20050003404A1

GENERAL INFORMATION:
TITLE OF INVENTION: TELEOMERASE INTERFERENCE
FILE REFERENCE: A-71506-1/RRT/THR

CURRENT APPLICATION NUMBER: US/10/831,266

CURRENT APPLICATION NUMBER: US/10/831,266

PRIOR APPLICATION NUMBER: US/00333065

PRIOR FILING DATE: 2002-10-16

PRIOR FILING DATE: 2001-10-22

PRIOR APPLICATION NUMBER: US 60/345,326

PRIOR FILING DATE: 2001-10-22

PRIOR APPLICATION NUMBER: US 60/345,326

PRIOR FILING DATE: 2001-10-22

PRIOR PILING DATE: 2002-0-20

PRIOR FILING DATE: 2002-0-20

PRIOR FILING DATE: 2002-0-22

NUMBER OF SEQ ID NOS: 17

SOFTWARE: PARCH NOS: 17

LENGTH: 21
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US-10-831-26-8

Sequence 8, Application US/10831266

Publication No. US200500034041

GENERAL INPORMATION:
APPLICANT: ROWIEY, PETER T.
TITLE OF INVENTION: TELOMERASE INTERFERENCE
FILE REFERENCE: A-71566-1/RFT/THR
CURRENT FILING DATE: 2004-04-22

PRIOR APPLICATION NUMBER: US/10/831,266

CURRENT FILING DATE: 2002-10-16

PRIOR APPLICATION NUMBER: US 60/345,326

PRIOR APPLICATION NUMBER: US 60/345,326

PRIOR PILING DATE: 2002-02-20

PRIOR PILING DATE: 2002-05-20

NUMBER OF SEQ ID NOS: 17
Query Match 4.2%; Score 19; DB Best Local Similarity 68.4%; Pred. No. 1e+0 Matches 13; Conservative 6; Mismatches
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                                                                                                            42 TTGTCTAACCCTAACTGAG 60
                                                                                                                                           1 UUGUCUAACCCUAACUGAG 19
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ORGANISM: Artificial
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LENGTH: 21
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Page 52

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; OTHER INFORMATION: oligonucleotide US-10-831-267-8
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                                                                                                                                                                            270 GGCTTCTCCGGAGGCACCC 288
                                                                                                                                                                                                         1 GGCTTCTCGGAGGCACCC 19
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US-10-923-330-546
                                                                                     Query Match
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                                                 Score 19; DB 1; Length 21;
Pred. No. 1e+02;
6; Mismatches 0; Indels
                                                                                              0; Indels
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                                                                                                                                                                                                                                                                                  Sequence 7, Application US/10811267
Sedence 7, Application US/10811267
GENERAL INPORMATION:
APPLICANT: ROALEY, Peter T.
TITLE OF INVENTION: TELOMERASE INTERFERENCE
FILE REFERENCE: A-7156-2/RFT/THR
CURRENT FILING DATE: 2004-04-22
PRIOR APPLICATION NUMBER: US/10/831,267
CURRENT FILING DATE: 2002-10-16
PRIOR APPLICATION NUMBER: US 60/345,326
PRIOR FILING DATE: 2001-10-22
PRIOR PELING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: US 60/359,196
PRIOR FILING DATE: 2002-02-0
PRIOR PELING DATE: 2002-02-0
PRIOR APPLICATION NUMBER: US 60/389,195
PRIOR PILING DATE: 2002-05-2
PRIOR PELING DATE: 2002-05-2
PRIOR PELING DATE: 2002-05-2
SEQ ID NOS: 23
SEQ ID NO 7
LENGTH: 21
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Sequence 8, Application US/10831267

Sequence 8, Application US/10831267

GENERAL INFORMATION:
TITLE OF INVENTION: TELCOMERASE INTERFERENCE
FILE REFERENCE: A-71506-2/RFT/THR
CURRENT APPLICATION NUMBER: US/10/831,267

CURRENT APPLICATION NUMBER: US/00/33146

PRIOR APPLICATION NUMBER: PCT/US 02/33146

PRIOR FILING DATE: 2001-10-16

PRIOR FILING DATE: 2001-10-16

PRIOR APPLICATION NUMBER: US 60/345,326

PRIOR FILING DATE: 2001-10-2

PRIOR APPLICATION NUMBER: US 60/359,196

PRIOR APPLICATION NUMBER: US 60/359,196

PRIOR APPLICATION NUMBER: US 60/383,195

PRIOR FILING DATE: 2002-02-2

PRIOR APPLICATION NUMBER: US 60/383,195

PRIOR APPLICATION NUMBER: US 60/383,195

PRIOR FILING DATE: 2002-05-22

NUMBER OF SEQ ID NOS: 23

SEQ ID NO 8

LENGTH:: 21
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                                                                                                                                                              1 UNGUCUAACCCUAACUGAG 19
                                                                                                                                    42 TTGTCTAACCCTAACTGAG 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         42 TIGICIAACCCIAACTGAG 60
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Best Local Similarity 68.4%;
Matches 13; Conservative (
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Best Local Similarity
Matches 19; Conserva
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ORGANISM: Artificial
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ORGANISM: Artificial
FEATURE:
                                                                                                                                                                                                                                                                    US-10-831-267-7/c
US-10-831-267-6
                                                                                          Matches
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     Length 21;
                                 0; Indels
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4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches 0; Indels
   DB 1;
1e+02;
                                                                                                                                                                  Sequence 9, Application US/10831267;
Sequence 9, Application US/10831267;
Fublication No. US20050009177A1
GENERAL INFORMATION:
TITLE OF INVENTION: TELOMERASE INTERFERENCE
FILE REFERENCE: A-71506-2/RFT/THR
CURRENT APPLICATION NUMBER: US/10/831,267
CURRENT FILING DATE: 2004-04-22
PRIOR APPLICATION NUMBER: US/00/345,326
PRIOR FILING DATE: 2001-10-16
PRIOR APPLICATION NUMBER: US 60/345,326
PRIOR FILING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: US 60/359,196
PRIOR PILING DATE: 2002-02-0
PRIOR PRICATION NUMBER: US 60/383,195
PRIOR FILING DATE: 2002-05-2
NUMBER OF SEQ ID NOS: 23
SOFTWARE: PALENTIN UN VERSION 3.2
SEQ ID NO 9
LENGTH: 21
Query Match
4.2%; Score 19; DB
Best Local Similarity 100.0%; Pred. No. 10+
Matches 19; Conservative 0; Mismatches
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FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: sinA sense region
                                                                         FEATURE:
NAME/KEY: misc_feature
LOCATION: (20)...(21)
OTHER INFORMATION: n stands for thymidine
US-10-923-330-547
                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 63.2%;
Matches 12; Conservative
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           g
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JEGUREAL INFORMATION:

JEGUREAL SEPTICANT:

JEGUREAL SEPTICANT:

JEGUREAL SEPTICANT:

JEGUREAL SEPTICANT:

JEGUREAL SEPTICANTION:

JEGUREAL SEPTICANT:

JEGUREAL SEPTICANTION NUMBER:

JEGUREAL SEPTICANTION SEPT
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OTHER INFORMATION: Description of Artificial Sequence: sinA sense region
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PRIOR PILING DATE: 2003-11-24
PRIOR APPLICATION NUMBER: US 10/693,059
PRIOR PILING DATE: 2003-11-23
PRIOR PLILING DATE: 2003-05-13
PRIOR APPLICATION NUMBER: PCT/US03/05346
PRIOR PILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR PILING DATE: 2003-02-20
PRIOR PILING DATE: 2003-02-20
REMAINING PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR PILING DATE: 2003-02-20
REMAINING PRIOR APPLICATION NUMBER: PCT/US03/05028
SOFTWARE: PATENTIN VERSION 3.3
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// LOCATION: (20)...(21)

OTHER INFORMATION: n stands for thymidine

US-10-923-330-546
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: RNA
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 546
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GENERAL INPOGNATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Beigen, James
APPLICANT: Beigen, Leonid
TITLE De INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE De INVENTION: Expression Using Short Interfering RNA (sinA)
FILE REFERENCE: 400/209
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT APPLICATION NUMBER: PCT/US03/04088
RRIOR APPLICATION NUMBER: PCT/US03/04088
RRIOR PILING DATE: 2004-05-24
FRIOR PILING DATE: 2004-05-24
FRIOR PILING DATE: 2004-01-14
FRIOR PILING DATE: 2003-01-1-3
FRIOR PILING DATE: 2003-01-1-3
FRIOR PILING DATE: 2003-01-23
FRIOR PILING DATE: 2003-05-23
FRIOR PILING DATE: 2003-05-23
FRIOR PILING DATE: 2003-05-24
FRIOR PILING DATE: 2003-05-20
FRIOR FILING DATE: 2003-05-20
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                                                                                  Gaps
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4.2%; Score 19; DB 1;
Best Local Similarity 68.4%; Pred, No. 1e+02;
Matches 13; Conservative 6; Mismatches
Score 19; DB 1;
Pred. No. 1e+02;
7; Mismatches
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NAME/KEY: misc feature
LOCATION: (20)...(21)
OTHER INFORMATION: n stands for thymidine
US-10-923-330-548
                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 548, Application US/10923330 Publication No. US20050153916A1
                                                                                                                                                                146 TICCACCGTICATICIAGA 164
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                         Sequence 549, Application US/10923330
| Publication No. US20050153916A1
| Publication No. US20050153916A1
| GENERAL INFORMATION:
| APPLICANT: Sirna Therapeutics, Inc.
| APPLICANT: Missigen, James
| TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
| FILE REFERENCE: 400/209 (WHENDE: US/10/923,330
| CURRENT FILING DATE: 2004-08-24
| PRIOR APPLICATION NUMBER: US/05-24
| PRIOR APPLICATION NUMBER: US 10/720,448
| PRIOR PELING DATE: 2004-01-14
| PRIOR PELING DATE: 2004-01-14
| PRIOR PELING DATE: 2003-11-24
| PRIOR PELING DATE: 2003-11-24
| PRIOR APPLICATION NUMBER: US 10/720,448
| PRIOR PELING DATE: 2003-05-23
| PRIOR PELING DATE: 2003-05-24
| PRIOR PELICATION NUMBER: US 10/444,853
| PRIOR PELICATION NUMBER: US PCT/US03/05026
| PRIOR PELICATION NUMBER: US PCT/US03/05026
| PRIOR PELICATION NUMBER: PCT/US03/05026
| PRIOR PELICATION NUMBER: PCT/US03/05026
| PRIOR PELICATION NUMBER: US PCT/US03/05026
| PRIOR PELICATION NUMBER: PCT/US03/05026
| PRIOR PELICATION PUMBER: PCT/US03/05026
| PCT/US03/
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APPLICANT: Sirma Therapeutics, Inc.
APPLICANT: Beigelnan, James
APPLICANT: Beigelnan, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Jahr Short Interfering RNA (siNA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: US/10/923, 330
CURRENT FILING DATE: 2004-08-20
PRIOR PILING DATE: 2004-08-24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: misc_feature

// LOCATION: (20)...(21)

US-10-923-330-549
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  148 CCACCGTTCATTCTAGAGC 166
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: RNA
ORGANISM: Artificial Sequence
US-10-923-330-549
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US-10-923-330-550
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HESULT 142

US-10-923-930-551

Sequence 551, Application US/10923330

Publication No. US20050153916A1

GENERAL INFORMATION:

APPLICANT: SITE THE THEORY JEAN

APPLICANT: MESWAGGEN, James

APPLICANT: MESWAGGEN, James

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

FILE REFERENCE: 400/209 (MHHRD2-708-C)

CURRENT APPLICATION NUMBER: 105/10/923,330

CURRENT APPLICATION NUMBER: D5/10/923,330

FRIOR PELING DATE: 2004-08-20

PRIOR APPLICATION NUMBER: D5/10/923,330

PRIOR APPLICATION NUMBER: D5/10/923,330

PRIOR APPLICATION NUMBER: D5/10/923,330

PRIOR PLING DATE: 2004-08-20

PRIOR APPLICATION NUMBER: US 60/396,600

PRIOR PLING DATE: 2004-01-14

PRIOR APPLICATION NUMBER: US 10/750,448

PRIOR APPLICATION NUMBER: US 10/750,448

PRIOR PLING DATE: 2004-01-14

PRIOR PLING DATE: 2003-11-24

PRIOR PLING DATE: 2003-11-24

PRIOR PLING DATE: 2003-11-23

PRIOR PLING DATE: 2003-11-23
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                                                                         FRIOR FILING DATE: 2002-07-17

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-04-16

PRIOR FILING DATE: 2004-04-16

PRIOR PELING DATE: 2004-04-16

PRIOR APPLICATION NUMBER: US 10/757,803

PRIOR PELING DATE: 2003-11-24

PRIOR PELING DATE: 2003-11-23

PRIOR PELING DATE: 2003-11-23

PRIOR PELING DATE: 2003-11-23

PRIOR PELING DATE: 2003-11-23

PRIOR PELING DATE: 2003-12-24

PRIOR APPLICATION NUMBER: PCT/US03/05346

PRIOR APPLICATION NUMBER: PCT/US03/05246

PRIOR PELING DATE: 2003-02-20

PRIOR PELING DATE: 2003-02-20

PRIOR PELING DATE: 2003-02-20

PRIOR SPLING DATE: 2003-02-20

PRIOR PELING DATE: 2003-02-20

PRIOR SPLING DATE: 2008-62-30

PRIOR PELING DATE: 2008-62-30

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: misc_feature; LOCATION: (20)...(21)
D. COCATION: (20)...(21)
US-10-923-330-550
APPLICATION NUMBER: US 60/396,600
FILING DATE: 2002-07-17
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ORGANISM: Artificial Sequence
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                                                                             - See File Wrapper or PALM.
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Pred. No. 1e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                   ; NAME/KEY: misc_feature
; LOCATION: (20)..(21)
; OTHER INFORMATION: n stands for thymidine
US-10-923-330-551
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; NAME/KEY: misc feature
; LOCATION: (20)...(21)
LOCATION: (20)...(21)
US-10-923-330-552
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR FILING DATE: 2003-02-20
Remaining Prior Application data removed
NUMBER OF SEQ ID NOS: 768
SOFTWARE: Patentin version 3.3
SEQ ID NO 551
LENGTH: 21
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ORGANISM: Artificial Sequence
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Best Local Similarity 73.7%;
Matches 14; Conservative
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US-10-923-330-552
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OTHER INFORMATION: Description of Artificial Sequence: BINA antisense region
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400/203 (MeHBD2-708-2)
CURRENT APPLICATION NUMBER: PCT/US03/04088
PRIOR PRILING DATE: 2004-605-24
PRIOR PELICATION NUMBER: US 60/396,600
PRIOR FILING DATE: 2004-04-14
PRIOR PELICATION NUMBER: US 10/926,966
PRIOR PELICATION NUMBER: US 10/757,803
PRIOR PELING DATE: 2004-04-14
PRIOR PELING DATE: 2004-01-14
PRIOR PELING DATE: 2003-11-24
PRIOR PELING DATE: 2003-01-27
PRIOR PELING DATE: 2003-05-23
PRIOR PELING DATE: 2003-05-23
PRIOR PELING DATE: 2003-05-23
PRIOR PELING DATE: 2003-02-20
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7. le+02;
0;
   DB 1;
                                                             5; Mismatches
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Score 19;
Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                 Sequence 554, Application US/10923330 Publication No. US20050153916A1
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                                                                                                                        300 GAAGAGTIGGGCTCTGTCA 318
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ORGANISM: Artificial Sequence
   Query Match
Best Local Similarity 73.7%;
Matches 14; Conservative
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US-10-923-330-555/c
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PRIOR APPLICATION NUMBER: 10 10/720,448
PRIOR FILING DATE: 2003-11-24
PRIOR FILING DATE: 2003-11-24
PRIOR FILING DATE: 2003-11-23
PRIOR FILING DATE: 2003-11-23
PRIOR FILING DATE: 2003-11-24
PRIOR PELING DATE: 2003-05-23
PRIOR PELING DATE: 2003-05-23
PRIOR APPLICATION NUMBER: PCT/US03/05346
PRIOR FILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR FILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR FILING DATE: 2003-02-20
PRIOR FILING DATE: 2003-02-20
PROBAINING PATE: 2003-02-20
PATE: 2003-
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ORGANISM: Artificial Sequence
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APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Rxpression Using Short Interfering RNA (siNA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-08-20
PRIOR FILING DATE: 2004-05-24
PRIOR PLING DATE: 2004-01-14
PRIOR PLING DATE: 2004-01-14
PRIOR PLING DATE: 2003-11-24
PRIOR PLING DATE: 2003-02-20
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Sequence 556, Application US/10923330

Publication No. US20050153916A1

GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400/209 (MRHBDZ-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330

CURRENT APPLICATION NUMBER: DC1/US03/04088

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2002-07-17

PRIOR APPLICATION NUMBER: PCT/US04/16390

PRIOR APPLICATION NUMBER: PCT/US04/16390

PRIOR PLING DATE: 2004-05-24

PRIOR PLING DATE: 2004-06-20

PRIOR FILING DATE: 2004-06-24

PRIOR PLING DATE: 2004-06-24

PRIOR PLING DATE: 2004-06-24

PRIOR PLING DATE: 2004-06-26

PRIOR PLING DATE: 2004-06-26

PRIOR PLING DATE: 2004-06-26

PRIOR PLING DATE: 2004-06-26

PRIOR APPLICATION NUMBER: US 10/826,966

PRIOR APPLICATION NUMBER: US 10/757,803
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NUMBER OF SEQ ID NOS: 768
SOFTWARE: PatentIn version 3.3
SEQ ID NO 555
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; LOCATION: (20)...(21)
... OTHER INFORMATION: n stands for thymidine
US-10-923-330-555
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ORGANISM: Artificial Sequence
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OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
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                                                            Gaps
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0
                                              Length 21;
                                                           0; Indels
                                              DB 1;
1e+02;
                                            Query Match 4.2%; Score 19; DB Best Local Similarity 100.0%; Pred. No. 1e-Matches 19; Conservative 0; Mismatches
     FEATURE:
NAME/KEY: misc feature
LOCATION: (20) ... (21)
COTHER INFORMATION: n stands for thymidine
US-10-923-330-556
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19 CACCGTTCATTCTAGAGCA 1
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APPLICANT: SITMS Therapellities, Inc.
APPLICANT: SITMS Therapellities, Inc.
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA INTERFERENCE: 400/209 (MBHB02-708-C)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT PAPLICATION NUMBER: US/10/923,330
CURRENT PAPLICATION NUMBER: DCT/US03/0408
PRIOR APPLICATION NUMBER: DCT/US03/0408
PRIOR PILING DATE: 2002-07-17
PRIOR PILING DATE: 2003-07-16
PRIOR PILING DATE: 2004-05-24
PRIOR PILING DATE: 2004-04-16
PRIOR PILING DATE: 2004-04-16
PRIOR PILING DATE: 2003-11-23
PRIOR PILING DATE: 2003-11-23
PRIOR PILING DATE: 2003-11-23
PRIOR PILING DATE: 2003-11-23
PRIOR PILING DATE: 2003-05-23
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                                                                                        OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
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Pred. No. 1e+02;
0; Mismatches 0; Indels
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4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                    ch 4.2%; Score 19; DB 1 Similarity 100.0%; Pred. No. 1e-19; Conservative 0; Mismatches
                                                                                                                     PEATURE:
NAME/KEY: misc_feature
LOCATION: (20) ... (21)
OTHER INFORMATION: n stands for thymidine
US-10-923-330-557
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LOCATION: (20)...(21)

COTHER INFORMATION: n stands for thymidine

US-10-923-330-558
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                                                                                                                                                                                                                                                                                                                                                                                                                                148 CCACCGTTCATTCTAGAGC 166
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19 CCACCGTTCATTCTAGAGC 1
TYPE: RNA
ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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LENGTH: 21
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Best Local 8
                                                            FEATURE:
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Page 58

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Sequence 563, Application US/10923330

Publication No. US20050153916A1

GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REPERBNCE: 400/209 (MHBD2-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT APPLICATION NUMBER: US/10/923,330
FRIOR APPLICATION NUMBER: DCT/US03/04088
FRIOR APPLICATION NUMBER: DCT/US03/04088
FRIOR PRILING DATE: 2004-05-24
FRIOR PILING DATE: 2002-07-17
FRIOR APPLICATION NUMBER: PCT/US04/16390
FRIOR PILING DATE: 2004-06-24
FRIOR APPLICATION NUMBER: PCT/US04/16390
FRIOR APPLICATION NUMBER: US/10/9204
FRIOR APPLICATION NUMBER: US/24
FRIOR PILING DATE: 2004-06-24
FRIOR APPLICATION NUMBER: US/24
FRIOR PILING DATE: 2004-06-40-66-51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: Description of Artificial Sequence: sinA sense region
                PRIOR APPLICATION NUMBER: PCT/US03/05346
PRIOR FILING DATE: 2003-02-20
PRIOR FILING DATE: 2003-02-20
PRIOR FILING DATE: 2003-02-20
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 768
SOFTWARE: Patentin version 3.3
LENGTH: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: misc_feature
LOCATION: (1)...(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(5)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY: misc_feature

LOCATION: (21)...(21)...

OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-562
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NAME/KEY: misc_feature
LOCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
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NAME/KEY: misc_feature
LOCATION: (7)...(8)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
FEATURE:
NAME/KEY: misc_feature
LOCATION: (10)...(12)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
FEATURE:

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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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1 CUUCCACCGUUCAUUCUAG 19
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Best Local Similarity
Matches 12; Conserva
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| Bublication No. US20050153916A1
| GENERAL INFORMATION:
| APPLICANT: Sirna Therapeutics, Inc.
| APPLICANT: Sirna Therapeutics, Inc.
| APPLICANT: Beigedman, Leonid
| TILE OF INVENTION: Expression Using Short Interfering RNA (sinA)
| TILE OF INVENTION: Expression Using Short Interfering RNA (sinA)
| TILE OF INVENTION: Expression Using Short Interfering RNA (sinA)
| FILE REFERENCE: 400/209 (WHHB02-708-C)
| FILE REFERENCE: 400/209 (WHHB02-708-C)
| FILE REFERENCE: 2004-08-20
| FRIOR APPLICATION NUMBER: PCT/US03/04088
| FRIOR FILING DATE: 2004-05-24
| FRIOR FILING DATE: 2004-05-24
| FRIOR FILING DATE: 2004-06-24
| FRIOR FILING DATE: 2004-01-14
| FRIOR FILING DATE: 2004-01-14
| FRIOR FILING DATE: 2004-01-14
| FRIOR FILING DATE: 2003-01-14
| FRIOR FILING DATE: 2003-11-24
| FRIOR FILING DATE: 2003-11-23
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                                                                                                                FRIOR FILING DATE: 2000-07-17
FRIOR APPLICATION NUMBER: DCT/US04/16390
FRIOR FILING DATE: 2004-05-24
FRIOR APPLICATION NUMBER: US 10/826,966
FRIOR APPLICATION NUMBER: US 10/826,966
FRIOR FILING DATE: 2004-01-14
FRIOR FILING DATE: 2004-01-14
FRIOR FILING DATE: 2003-11-24
FRIOR FILING DATE: 2003-11-24
FRIOR FILING DATE: 2003-11-24
FRIOR PLILING DATE: 2003-11-24
FRIOR PLILING DATE: 2003-12-24
FRIOR PLILING DATE: 2003-12-24
FRIOR PLILING DATE: 2003-05-23
FRIOR PLILING DATE: 2003-05-23
FRIOR APPLICATION NUMBER: US 10/444,853
FRIOR APPLICATION NUMBER: PCT/US03/05346
FRIOR PLILING DATE: 2003-02-20
FRIOR FILING DATE: 2003-02-30
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APPLICATION NUMBER: PCT/US03/04088
                                                FILING DATE: 2004-05-24
APPLICATION NUMBER: US 60/396,600
FILING DATE: 2002-07-17
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US-10-923-330-565
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Publication No. US20050153916A1

GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Beigefinan, Leonid
TITLE OF INVENTION: RN Interference Mediated Inhibition of Telomerase General Engerance: 400,209 (MRH802-708-C)
FILE REPRENENCE: 400,209 (MRH802-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-08-20
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PRIOR APPLICATION NUMBER: US 10/757,803
PRIOR FILING DATE: 2004-01-14
PRIOR PLILING DATE: 2003-11-24
PRIOR PLILING DATE: 2003-11-23
PRIOR PLILING DATE: 2003-11-23
PRIOR PLILING DATE: 2003-01-12-3
PRIOR PLILING DATE: 2003-05-23
PRIOR PLILING DATE: 2003-05-20
PRIOR PLILING DATE: 2003-02-20
PRIOR PRIOR DATE: 2003-02-20
PRIOR PLILING DATE: 2003-02-20
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LOCATION: (1)._(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
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LOCATION: (21) ... (21)
OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
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4.2%; Score 19; DB 1;
Best Local Similarity 63.2%; Pred. No. 1e+02;
Matches 12; Conservative 7; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: misc_feature
LOCATION: (20) ... (20)
OTHER INFORMATION: n stands for thymidine
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LOCATION: (9)._(11)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (13)...(16)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (1)._(4)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (6)._(7)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: RNA
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sinA sense region
FRIOR APPLICATION UNDERS: 306/396,600
FRIOR FILING DATE: 2002-07-17
FRIOR PELING DATE: 2002-07-17
FRIOR PELING DATE: 2004-05-24
FRIOR FILING DATE: 2004-05-24
FRIOR FILING DATE: 2004-05-24
FRIOR FILING DATE: 2004-01-46
FRIOR PELING DATE: 2004-01-46
FRIOR PELING DATE: 2004-01-48
FRIOR PELING DATE: 2004-01-49
FRIOR PELING DATE: 2003-11-24
FRIOR PELING DATE: 2003-11-24
FRIOR PELING DATE: 2003-11-23
FRIOR PELING DATE: 2003-11-23
FRIOR PELING DATE: 2003-11-23
FRIOR PELING DATE: 2003-05-23
FRIOR PELING DATE: 2003-05-23
FRIOR PELING DATE: 2003-05-23
FRIOR PELING DATE: 2003-05-20
FRIOR FILING DATE: 2003-05-20
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OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
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OTHER INFORMATION: n stands for thymidine
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LOCATION: (1)._(3)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            147 TCCACCGTTCATTCTAGAG 165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 UCCACCGUUCAUUCUAGAG 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: RNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: misc feature
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LOCATION: (12)..(15)
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LOCATION: (20)..(20)
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Wed Aug 24 14:55:17 2005

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**PELLICANT: O-1111 AUG-SWIGGERULICE, THE.**

**APPLICANT: Beigelman, Leonid

**PELLICANT: Beigelman, Leonid

**PELLICANTON: RANA Interference Mediated Inhibition of Telomerase Gene

**PELLICANTON: TEXPERSION: TOWABER: US/10/923,330

**CURRENT APPLICATION NUMBER: US/10/923,330

**PELLING DATE: 2004-05-24

**PRIOR FILING DATE: 2002-07-17

**PRIOR FILING DATE: 2004-05-24

**PRIOR FILING DATE: 2004-05-24

**PRIOR PELLING DATE: 2004-06-24

**PRIOR FILING DATE: 2004-01-14

**PRIOR FILING DATE: 2004-01-14

**PRIOR FILING DATE: 2003-01-12

**PRIOR FILING DATE: 2003-11-24

**PRIOR PLING DATE: 2003-05-23

**PRIOR PLING DATE: 2003-05-23

**PRIOR PLING DATE: 2003-02-20

**PRIOR PLING DATE: 2003-02-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        siNA sense region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 768
SOFTWARE: Patentin version 3.3
SEQ ID NO 565
LENGTH: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   i LOCATION: (21)...(21)...
i OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity US-10-923-330-565
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Description of Artificial Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LOCATION: (20)..(20)
OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: misc feature
LOCATION: (1)._(2)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: misc feature
LOCATION: (4)._(5)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DOCATION: (/)..(9)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: misc_feature
LOCATION: (11)...(14)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: misc feature
LOCATION: (19)...(19)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
APPLICANT: Sirna Therapeutics, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: misc_feature
LOCATION: (7)..(9)
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NAME/KEY: misc feature
LOCATION: (20)..(20)
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LOCATION: (21)..(21)
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Length 21;

Score 19; DB 1; Pred. No. 1e+02;

4.2%;

Query Match Best Local Similarity

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WESULT 123-356-566

WESULT 125-1330-566

Sequence 566, Application US/10923330

GENERAL INFORMATION:
APPLICANT: SITEM Therapeutics, Inc.
APPLICANT: HIGH Therapeutics, Inc.
APPLICANT: Helgelman, Leonid
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
CURRENT APPLICATION NUMBER: US (004-08-20)
PRIOR PELICAN ADMER: 2004-07-17
PRIOR PELICAN UNMER: US 10/826,966
PRIOR PELICANION NUMBER: US 10/826,966
PRIOR PELICANION NUMBER: US 10/720,448
PRIOR PELICANION NUMBER: US 10/720,448
PRIOR APPLICATION NUMBER: US 10/720,448
PRIOR APPLICATION NUMBER: US 10/720,448
PRIOR APPLICATION NUMBER: US 10/303/05346
PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR APPLICATION NUMBER: PCT/US03/0528
PRIOR PRIOR PLING DATE: 2003-02-20
PRIOR PRIOR PLING DATE: 2003-02-20
PRIOR PRIOR PLING DATE: 2003-02-20
PRIOR PLING DATE: 
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           Gaps
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LOCATION: (1)._(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
              Indels
           .;
0
       5; Mismatches
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NAME/KEY: misc_feature
LOCATION: (1)._(1)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (3).7(4)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY: misc_feature
LOCATION: (10)..(13)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (18)...(18)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                              TYPE: RNA
ORGANISM: Artificial Sequence
   Conservative
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LOCATION: (6). (8)
14;
   Matches
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19-10-923-330-56

19-10-923-330-56

19-10-10-923-330-56

19-10-10-92-10-926

19-10-10-92-10-926

19-10-10-92-10-926

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NAME/KEY: misc_feature
LOCATION: (1)._(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
                                                                                                                                                                                    NAME/KEY: misc_feature

1 LOCATION: (21)...(21)

2 OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity

US-10-923-330-566
                                                                                                                                                                                                                                                                                                                                                                 Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 73.7%; Pred. No. 1e+02; Matches 14; Conservative 5; Mismatches 0; Indels
FEATURE:
NAME/KEY: misc_feature
LOCATION: (20)..(20)
OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: misc_feature
LOCATION: (5)._(7)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
MAME/KEX: misc_feature
LOCATION: (2). (3)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           149 CACCGTTCATTCTAGAGCA 167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: RNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 156
US-10-923-330-567
                                                                                                                                                     FEATURE:
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## STATUS APPLICANT NO. USZO050153916A1

| GENERAL INFORMATION:
| APPLICANT: Sirna Therapeutics, Inc.
| APPLICANT: Sirna Therapeutics, Inc.
| APPLICANT: Mosaiggen, James
| APPLICANT: Mosaiggen, James
| APPLICANT: Baigelman, Leonid
| TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
| TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
| TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
| TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
| TITLE OF INVENTION: RNA INTERF. US/10/923,330
| CURRENT FILING DATE: 2004-08-24
| PRIOR FILING DATE: 2004-06-24
| PRIOR FILING DATE: 2004-01-16
| PRIOR FILING DATE: 2004-01-16
| PRIOR APPLICATION NUMBER: US 10/757,803
| PRIOR PELICATION NUMBER: US 10/757,803
| PRIOR FILING DATE: 2004-01-14
| PRIOR FILING DATE: 2004-01-14
| PRIOR FILING DATE: 2003-11-24
| PRIOR FILING DATE: 2003-11-24
| PRIOR FILING DATE: 2003-11-23
| PRIOR PAPLICATION NUMBER: US 10/693,059
| PRIOR FILING DATE: 2003-02-10
| PRIOR PILING DATE: 2003-02-20
| PRIOR PILING DATE: 2003-02-30
| PRIOR PILING D
                                                                                                                                                                                                                                                                                                                                                                                                                                                 LOCATION: (20)...(20) OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
4.2%; Score 19; DB 1;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; LOCATION: (21) ... (21)
; OTHER INFORMATION: n stands for thymidine
US-10-923-330-570
                                                                                                                NAME/KEY: misc feature
LOCATION: (7)...(7)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                     DOCATION: (1)..(2)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                   NAME/KEY: misc feature
LOCATION: (11)...(11)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                   NAME/KEY: misc feature
LOCATION: (14)...(14)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 571, Application US/10923330
Publication No. US20050153916A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            145 CTTCCACCGTTCATTCTAG 163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19 CTTCCACCGTTCATTCTAG 1
                     misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
NAME/KEY: misc feature
LOCATION: (21)...(21)
                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: misc feature
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                                                                                                                                                                                        FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
NAME/KEY: misc_feature
LOCATION: (21)...(21)
COTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-568
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 73.7%; Pred. No. 10+02;
Matches 14; Conservative 5; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                   LOCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
                                                                                                NAME/KEY: misc feature
LOCATION: (12)...(15)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
FEATURE:
NAME/KEY: misc feature
LOCATION: (17)...(18)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
NAME/KEY: misc feature
LOCATION: (7)..(8)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                        NAME/KEY: misc_feature
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Gaps

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Length 21; 0; Indels

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Sequence 573, Application US/10923330

Publication No. US20050153916A1

GENERAL INPORMATION:

APPLICANT: Birna Therapeutics, Inc.
APPLICANT: Belgelman, Leonid

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

FILE REFERENCE: 400/209 (MRHBO2-708-C)

CURRENT FILING DATE: 2004-08-20

FRIOR PEPLICATION NUMBER: PCT/US03/0408B

PRIOR PELING DATE: 2004-05-24

PRIOR PELING DATE: 2004-01-14

PRIOR PELING DATE: 2004-01-14

PRIOR PELING DATE: 2004-01-14

PRIOR PELING DATE: 2003-11-24

PRIOR PELING DATE: 2003-11-24

PRIOR PELING DATE: 2003-11-23

PRIOR PELING DATE: 2003-11-23

PRIOR PELING DATE: 2003-11-23

PRIOR PELING DATE: 2003-11-23

PRIOR PELING DATE: 2003-05-23

PRIOR PELING DATE: 2003-05-23
                                                                                                                                                                                                                         INFORMATION: Description of Artificial Sequence: sinh antisense region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 100.0%; Pred. No. 1e+02; Matches 19; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; LOCATION: (21)...(21); OTHER INFORMATION: n stands for thymidine US-10-923-330-572
                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: misc_feature
LOCATION: (9)._(9)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
NAME/KEY: misc_feature
LOCATION: (16)..(16)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                    NAME/KEY: misc_feature
LOCATION: (1)...(4)
OCHER INFORMATION: 2'-deoxy-2'-fluoro
FEATURE:
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NUMBER OF SEQ ID NOS: 768
SOFWRARE: Patentin version 3.3
SEQ ID NO 572
LENGTH: 21
                                                                                                                                                              ORGANISM: Artificial Sequence
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NAME/KEY: misc_feature
LOCATION: (21)...(21)
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US-10-923-330-573/c
                                                                                                                                                                                                                                                             FEATURE:
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APPLICANT: Birna Therapeutics, Inc.
APPLICANT: Birna Therapeutics, Inc.
APPLICANT: Baggelann. James
APPLICANT: Baggelann. James
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-05-24
PRIOR PELIOR DATE: 2004-05-24
PRIOR APPLICATION NUMBER: US 60/396,600
PRIOR APPLICATION NUMBER: US 10/826,966
PRIOR FILING DATE: 2004-04-16
PRIOR PELING DATE: 2004-01-14
PRIOR PELING DATE: 2004-01-14
PRIOR PELING DATE: 2003-11-24
PRIOR PELING DATE: 2003-11-24
PRIOR PELING DATE: 2003-11-23
PRIOR PELING DATE: 2003-11-23
PRIOR APPLICATION NUMBER: US 10/757,803
PRIOR PELING DATE: 2003-11-23
PRIOR PELING DATE: 2003-11-24
PRIOR PELING DATE: 2003-10-220
PRIOR PELING DATE: 2003-10-220
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR PELING DATE: 2003-02-20
                                                                                                  OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
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LOCATION: (20)...(20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LOCATION: (21)...(21)
COTHER INFORMATION: n stands for thymidine US-10-923-330-571
                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: misc feature
LOCATION: (12)...(12)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY: misc_feature
LOCATION: (15)..(15)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)..(3)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME:
NAME: misc_feature
LCATION: (8). (8). (8)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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; Sequence 572, Application US/10923330
; Publication No. US20050153916A1
; GENERAL INFORMATION:
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                                    ORGANISM: Artificial Sequence
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                                                                        FEATURE
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NAME/KEY: misc_feature

: LOCATION: (21)...(21)

: OTHER INFORMATION: n stands for thymidine

US-10-923-330-574
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NAME/KEY:
LOCATION: (11). (11)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
FEATURE:
NAME/KEY: misc feature
LOCATION: (15). (15)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: misc feature
LOCATION: (1)._(1)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
NAME/KEY: misc_feature
LOCATION: (3)..(6)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE.
NAME/KEY: misc_feature
LOCATION: (18)..(18)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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                                                                                                                                                                                                                                                         SEQ ID NO 574
                                                                                                                                                                                                                                                                                                        TYPE: RNA
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Publication No. US20050153916A1

GENERAL INFORMATION:

APPLICANT: Birna Therapeutics, Inc.

APPLICANT: McSwiggen, James

APPLICANT: McSwiggen, James

APPLICANT: Begelman, Leonid

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

CURRENT APPLICATION NUMBER: US/10/923,330

CURRENT PILING DATE: 2004-06-20

PRIOR PILING DATE: 2004-05-20

PRIOR APPLICATION NUMBER: US 60/396,600

PRIOR PILING DATE: 2004-05-24

PRIOR PILING DATE: 2004-05-24

PRIOR PLILNG DATE: 2004-06-16

PRIOR FILING DATE: 2004-06-16

PRIOR FILING DATE: 2004-06-16

PRIOR FILING DATE: 2004-01-14

PRIOR FILING DATE: 2004-01-14

PRIOR FILING DATE: 2004-01-14

PRIOR FILING DATE: 2004-01-14

PRIOR FILING DATE: 2003-11-24

PRIOR APPLICATION NUMBER: US 10/593,059
                                                                                                                                                                                                                              FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
FEATURE:
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PRIOR FILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR FILING DATE: 2003-02-20
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 768
SOFTWARE: Patentin version 3.3
LENGTH: 21
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LOCATION: (20)...(20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 19; DB 1; Length 21; Pred. No. 1e+02; 0; Mismatches 0; Indels
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100.0%; Pred. No. 1c...
... 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; LOCATION: (21)...(21)
; OTHER INFORMATION: n stands for thymidine
US-10-923-330-573
                                                                                                                                                                                                                                                                                                           NAME/KEY: misc_feature
LOCATION: (2)...(5)
OTHER_INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (14)...(14)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (17) ... (17)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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                                                                                                                                                                                                             ORGANISM: Artificial Sequence
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Best Local Similarity 100.
Matches 19; Conservative
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NAME/KEY: misc feature
LOCATION: (10)...(10)
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NAME/KEY: misc_feature
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                                                                                                                                                                                              TYPE: RNA
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Sequence 576, Application US/10923330

Publication No. US20050153916A1

APPLICANT: Britan Therapeutics, Inc.

APPLICANT: McSwiggen, James

APPLICANT: Beigelman, Leonid

TITLE OF INVENTION: RAN INTERference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

TITLE OF INVENTION: RANDER: US/10/203,330

CURRENT APPLICATION NUMBER: US/10/203,330

CURRENT FILING DATE: 2004-08-20

PRIOR FILING DATE: 2004-05-24

PRIOR APPLICATION NUMBER: US 60/396,600

PRIOR FILING DATE: 2002-07-17

PRIOR PRILING DATE: 2002-07-17

PRIOR APPLICATION NUMBER: PCT/US04/16390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
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PRIOR FILING DATE: 2003-11-23
PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR FILING DATE: 2003-05-23
PRIOR PLIING DATE: 2003-02-20
PRIOR PILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR FILING DATE: 2003-02-20
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 768
SOFTWARE: Patentin version 3.3
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NAME/KEY: misc_feature
LOCATION: (20)..(20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
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4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches 0; Indels
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Sequence 578, Application US/10923330; Publication No. US20050153916A1; GENERAL INFORMATION: APPLICANT: Sirna Therapeutics, Inc.
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NAME/KEY: misc_feature
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NAME/KEY: misc_feature
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NAME/KEY: misc_feature
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Sequence 576, Application US/10923330

Publication No. US20050153916A1

GENERAL INFORMATION:

APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: McSwiggen, James

APPLICANT: Beigelman, Leonid

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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LOCATION: (20) ... (20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 19; DB 1; Length 21; Pred. No. 1e+02; 0; Mismatches 0; Indels
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100.0%; Pred. No. 100.0%; O; Mismatches
PRIOR FILING DATE: 2004-05-24
PRIOR APPLICATION NUMBER: US 10/826,966
PRIOR PILING DATE: 2004-04-16
PRIOR PELICATION NUMBER: US 10/757,803
PRIOR FILING DATE: 2004-01-14
PRIOR PILING DATE: 2004-01-14
PRIOR FILING DATE: 2003-11-23
PRIOR FILING DATE: 2003-11-23
PRIOR FILING DATE: 2003-11-23
PRIOR FILING DATE: 2003-05-23
PRIOR FILING DATE: 2003-05-23
PRIOR FILING DATE: 2003-05-23
PRIOR FILING DATE: 2003-05-20
PRIOR PILING DATE: 2003-05-20
PRIOR FILING DATE: 2003-05-20
PRIOR FILING DATE: 2003-02-20
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j OTHER INFORMATION: n stands for thymidine
US-10-923-330-575
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
NAME/KEY: misc_feature
LOCATION: (4)..(7)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
NAME/KEX: misc_feature
LOCATION: (19)..(19)
OTHER_INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KRY: misc_feature
LOCATION: (1). (2)
OTHER INPORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: RNA
ORGANISM: Artificial Sequence
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Best Local Similarity 100.0
Matches 19; Conservative
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NAME/KEY: misc_feature
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OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
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CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT APPLICATION NUMBER: US/10/923,330
PRIOR APPLICATION NUMBER: PCT/US03/04088
PRIOR PILING DATE: 2004-06-20
PRIOR FILING DATE: 2004-05-24
PRIOR PILING DATE: 2004-05-24
PRIOR PILING DATE: 2004-05-24
PRIOR PILING DATE: 2004-04-64
PRIOR PILING DATE: 2004-04-64
PRIOR PILING DATE: 2004-04-64
PRIOR PILING DATE: 2004-04-14
PRIOR PILING DATE: 2003-11-24
PRIOR PILING DATE: 2003-12-23
PRIOR PILING DATE: 2003-05-23
PRIOR PILING DATE: 2003-05-23
PRIOR PILING DATE: 2003-02-20
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LOCATION: (20) ... (20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 21;
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4.2%; Score 19; DB 1; Le:
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: n stands for thymidine US-10-923-330-576
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LOCATION: (4)...(4)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY: misc_feature
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APPLICANT: Bejeglman, leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT PILING DATE: 2004-08-20
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-24
PRIOR PRIOR APPLICATION NUMBER: US 60/396,600
PRIOR PRILING DATE: 2004-05-24
PRIOR APPLICATION NUMBER: US 10/826,966
PRIOR PILING DATE: 2004-05-24
PRIOR PELING DATE: 2004-01-16
PRIOR PELING DATE: 2003-11-24
PRIOR PELING DATE: 2003-11-24
PRIOR PELING DATE: 2003-11-23
PRIOR PELING DATE: 2003-11-23
PRIOR PELING DATE: 2003-11-23
PRIOR PELING DATE: 2003-05-23
PRIOR PELING DATE: 2003-05-23
PRIOR PELING DATE: 2003-05-20
PRIOR FILING DATE: 2003-05-20
PRIOR PELING DATE: 2003-05-20
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LOCATION: (1)._(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
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LOCATION: (1)...(5)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (7)...(8)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY: misc_feature
LOCATION: (14)..(17)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (10)..(12)
TYTER INFORMATION: 2'-deoxy-2'-fluoro
FRATURE:
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ORGANISM: Artificial Sequence
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LOCATION: (6). (6)
OTHER INFORMATION: 2'-deoxy
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THER INFORMATION: 2'-deoxy
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LOCATION: (13)...(13)
OTHER INFORMATION: 2'-deoxy
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JOCATION: (18)..(19)
JTHER INFORMATION: 2'-deoxy
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NAME/KEY: misc_feature
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FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: sinA sense region
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR FILING DATE: 2003-02-20
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 76 NOS: 76 SEQ ID NOS SEQ ID NO SE
                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: misc feature
LOCATION: (1)..(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
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; OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-580
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Pred. No. 1e+02;
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OTHER INFORMATION: n stands for thymidine
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: misc feature
LOCATION: (1). (3)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: 2'-deoxy-2'-fluoro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   147 TCCACCGTTCATTCTAGAG 165
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                                                                                                                                                                                                                                                                  TYPE: RNA
ORGANISM: Artificial Sequence
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Best Local Similarity 68.4%;
Matches 13; Conservative
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LOCATION: (4). . . (4)
OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
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US-10-923-330-560

Sequence 580, Application US/10923330

Sequence 580, Application US/10923330

Sequence 580, Application US/10923330

Sequence 580, Application US/10923330

Sequence 580, Application of US/10923330

SEGUENCAL INFORMATION:
APPLICANT: Siring Therapeutics, Inc.
APPLICANT: Baigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: UNMERS: US/10/923,330

CURRENT APPLICATION WUMBER: US/10/923,330

CURRENT PILING DATE: 2004-08-20

PRIOR APPLICATION NUMBER: US 60/396,600

PRIOR FILING DATE: 2004-06-24

PRIOR APPLICATION NUMBER: US 10/326,600

PRIOR PILING DATE: 2004-01-14

PRIOR PILING DATE: 2004-01-14

PRIOR PLILING DATE: 2004-01-14

PRIOR PLILING DATE: 2003-11-23

PRIOR PLILING DATE: 2003-11-23

PRIOR PLILING DATE: 2003-11-23

PRIOR FILING DATE: 2003-02-20

PRIOR FILING DATE: 2003-02-20

PRIOR PLILING DATE: 2003-02-20

PRIOR FILING DATE: 2003-02-20
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; OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-579
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4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 63.2%; Pred. No. 1e+02;
Matches 12; Conservative 7; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: misc_feature
LOCATION: (20)_.(20)
OTHER INFORMATION: n stands for thymidine
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NAME/KEY: misc_feature
LOCATION: (9)..(11)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: misc feature
LOCATION: (13)...(16)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                     LOCATION: (6).7(7)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 UUCCACCGUUCAUUCUAGA 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LOCATION: (12)...(12)
OTHER INFORMATION: 2'-deoxy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LOCATION: (1/)..(15)
OTHER INFORMATION: 2'-deoxy
                                                                                                                                                                                             LOCATION: (8)..(8)
OTHER INFORMATION: 2'-deoxy
               NAME/KEY: misc feature
                                                                                                                     FEATURE:
NAME/KEY: misc_feature
LOCATION: (8). (8)
                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: misc feature
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Gaps

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Length 21; Indels

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LOCATION: (19)..(19)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                    NAME/KEY: misc_feature
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### APPLICANT: Beigelman, Leonid
### TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
### TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
### CURRENT APPLICATION NUMBER: US/10/923,330
### CURRENT PELICATION NUMBER: US 60/396,600
### PRIOR PELICATION NUMBER: US 60/396,600
### PRIOR PELING DATE: 2004-05-24
### PRIOR PELING DATE: 2004-06-24
### RIOR PELING DATE: 2004-01-16
### PRIOR PELING DATE: 2004-01-14
### PRIOR PELING DATE: 2004-01-14
### PRIOR PELING DATE: 2003-01-24
### PRIOR PELING DATE: 2003-01-24
### PRIOR PELING DATE: 2003-11-23
### PRIOR PELING DATE: 2003-11-23
### PRIOR PELING DATE: 2003-02-20
#### PRIOR PELING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Description of Artificial Sequence: sinA sense region FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: misc feature
LOCATION: (1). (1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(2)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (4)..(5)
OTHER_INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: misc feature
LOCATION: (7). (9)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (11)...(14)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: RNA
ORGANISM: Artificial Sequence
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LOCATION: (3)...(3)
OTHER INFORMATION: 2'-deoxy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
NAME/KEY: misc_feature
LOCATION: (6)...(6)
OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
FEATURE:
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NAME/KEY: misc_feature
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                                                                                                             Gaps
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                              NAME/KEY: misc_feature; LOCATION: (21)...(21); OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity US-10-923-330-581
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LOCATION: (1)..(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
                                                                                 Query Match
4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 73.7%; Pred. No. 1e+02;
Matches 14; Conservative 5; Mismatches 0; Indels
LOCATION: (20) ... (20)
OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
NAME/KEY: m.sc_feature
LOCATION: (1)...(1)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
FEATURE:
NAME/KEY: misc_feature
                                                                                                                           148 CCACCGTTCATTCTAGAGC 166
                                                                                                                                         1 CCACCGUUCAUUCUAGAGC 19
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ORGANISM: Artificial Sequence
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PRIOR FILING DATE: 2004-01-13
PRIOR PELING DATE: 2004-01-14
PRIOR PELING DATE: 2004-01-12
PRIOR PELING DATE: 2003-11-24
PRIOR PILING DATE: 2003-11-23
PRIOR PILING DATE: 2003-11-23
PRIOR PILING DATE: 2003-05-23
PRIOR PILING DATE: 2003-05-23
PRIOR PILING DATE: 2003-02-20
PRIOR PELING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR PILING DATE: 2003-02-20
PRIOR PILING DATE: 2003-02-20
PRIOR PILING DATE: 2003-02-20
PRIOR FILING DATE: 2003-02-30
PRIOR FILING DATE: 2003-03-30
PRIOR FILING DATE: 2003-03-30
PRIOR FILING DATE: 2003-03-30
PRIOR FILING PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR FILING DATE: 2003-03-30
PRIOR PILING DATE: 2003-03-30
PRIOR FILING DATE: 2003-03-33
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FILING DATE: 2004-04-16
APPLICATION NUMBER: US 10/757,803
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (17)...(17)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LOCATION: (5)...(7)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LOCATION: (9). 7(12)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: misc feature
LOCATION: (1)...(1)
OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
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LOCATION: (18)...(19)
OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
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LOCATION: (4)...(4)
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NAME/KEY: misc_feature
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LENGTH: 21
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US-10-923-330-583

US-10-923-330-583

Sequence 583, Application US/10923330

Publication No. US20050153916A1

GENERAL INFORMATION:

APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: Beigelman, Leonid

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RNA INTERFERENCE: 400/209 (WBHB02-708-C)

FILE REFERENCE: 400/209 (WBHB02-708-C)

CURRENT FILING DATE: 2004-08-20

PRIOR PLICATION NUMBER: PCT/US03/04088

PRIOR APPLICATION NUMBER: BCT/US03/04088

PRIOR PLING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24
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i LOCATION: (21)...(21)
i URPORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-582
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 73.7%; Pred. No. 1e+02; Matches 14; Conservative F. Wiemerband
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
NAME/KEY: misc feature
LOCATION: (20) (20)
OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: misc feature
LOCATION: (10) ... (13)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: misc feature
LOCATION: (18) ... (18)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                            FEATURE:
NAME/KEX: misc_feature
LOCATION: (6)..(8)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                       LOCATION: (3). 7(4)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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                                                                                                                                                     FEATURE:
NAME/KEY: misc_feature
LOCATION: (5)...(5)
OTHER INFORMATION: 2'-deoxy
                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: misc_feature
LOCATION: (9)..(9)
OTHER INFORMATION: 2'-deoxy
LOCATION: (2)..(2)

STHER INFORMATION: 2'-deoxy
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LOCATION: (14)...(17)
OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14; Conservative
                                                                            NAME/KEY: misc_feature
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OTHER INFORMATION: Description of Artificial Sequence: sinA sense region
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; OTHER INFORMATION: 3'-3 attached terminal deoxyabaaic moeity
US-10-923-330-583
                                                                                                                   LOCATION: (1). (1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
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OTHER INFORMATION: n stands for thymidine
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DB 1; Length 21;

US09436060A-16.rnpb.sl

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OTHER INFORMATION: 2'-deoxy
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9 Publication No. US20050153916A1
GENERAL INFORMATION:
1 APPLICANT: Sirna Therapeutics, Inc.
1 APPLICANT: McSwigen, James
APPLICANT: McSwigen, James
APPLICANT: Beigelman, Leonid
1 TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
1 TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
1 TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
1 FILE REPRENCE: 400/209 (MHHBO)-708-C)
1 CURRENT APPLICATION NUMBER: US/10/923,330
1 FILE MC APPLICATION NUMBER: US (00/30)/4088
1 FRIOR FILING DATE: 2004-08-20
1 FRIOR FILING DATE: 2002-07-17
1 FRIOR FILING DATE: 2004-01-4
1 FRIOR FILING DATE: 2004-01-14
1 FRIOR FILING DATE: 2003-11-23
1 FRIOR FILING DATE: 2003-01-24
1 FRIOR FILING DATE: 2003-02-20
1 FRIOR FILING DATE: 2003-02-20
2 FRIOR FILING DATE: 2003-02-20
3 FRIOR FILING DATE: 2003-02-20
3 FRIOR FILING DATE: 2003-02-20
3 FRIOR FILING DATE: 2003-02-20
4 FRIOR FILING DATE: 2003-02-20
5 FRIOR FILING DATE: 2003-03-20
5 FRIOR FILING DATE: 2003-03-30
5 FRIOR FILING DATE: 20
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                                                  Gaps
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LOCATION: (1)._(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
                                              0; Indels
73.7%; Pred. No. 1e+02;
tive 5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: misc feature
LOCATION: (7)...(8)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (12)...(15)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                           150 ACCGTTCATTCTAGAGCAA 168
                                                                                                                                                      TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: misc feature
LOCATION: (1). (6)
OTHER INFORMATION: 2'-deoxy
                                      14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
NAME/KEY: misc_feature
LOCATION: (9)._(11)
OTHER INFORMATION: 2'-deoxy
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: misc feature LOCATION: (16)..(16)
   Best Local Similarity
Matches 14; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 584
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Sequence 566, Application US/10923330

Sequence 566, Application US/10923330

Publication No. US20050153916A1

GENERAL INFORMATION:

APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: McSWagen, James

APPLICANTON WDARE: 2004-06-20

FRICR RILING DATE: 2004-06-24

FRICR PILING DATE: 2004-01-17

FRICR PILING DATE: 2004-01-16

FRICR PILING DATE: 2004-01-16

FRICR PILING DATE: 2004-01-16

FRICR PILING DATE: 2004-01-16

FRICR PILING DATE: 2003-01-12

FRICR FILING DATE: 2003-11-24

FRICR FILING DATE: 2003-11-24

FRICR FILING DATE: 2003-01-14

FRICR FILING DATE: 2003-01-14

FRICR FILING DATE: 2003-01-14

FRICR FILING DATE: 2003-02-20

FRICR FILING DA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                               NAME/KEY: misc_feature
) LOCATION: (20)
COTHER INFORMATION: n stands for thymidine
) FEATURE:
NAME/KEY: misc_feature
LOCATION: (21)
COTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-584
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 19; DB 1;
Pred. No. 1e+02;
5; Mismatches
NAME/KEY: misc feature
LOCATION: (17)...(18)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        300 GAAGAGTTGGGCTCTGTCA 318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 GAAGAGUUGGGCUCUGUCA 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 73.7%;
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: RNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                   NAMES/KEY: misc feature
LOCATION: (19)...(19)
OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
                  PRIOR FILING DATE: 2003-11-23
PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR FILING DATE: 2003-02-23
PRIOR FILING DATE: 2003-02-20
PRIOR FILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: PCT/US03/0508
PRIOR APPLICATION NUMBER: PCT/US03/05028
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 768
SOFTWARE: Patentin version 3.3
EEQ ID NO 587
LENGTH: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: misc_feature

COGATION: (20)...(20)

OTHER INFORMATION: Phosphorothicate 3'-Internucleotide Linkage
PEATURE:
NAME/KEY: misc_feature

LOCATION: (21)...(21)

COTHER INFORMATION: n stands for thymidine

US-10-923-330-587
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APPLICATION NUMBER: US 10/693,059
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LOCATION: (8). (8) OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: misc feature
LOCATION: (12)...(12)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LOCATION: (15)...(15)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LOCATION: (1)..(3)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: misc feature
LOCATION: (9)..(11)
OTHER INFORMATION: 2'-deoxy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LOCATION: (4). (7)
OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
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NAME/KEY: misc_feature
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LOCATION: (16)..(19)
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                                                                                                                                                                                                                                                                                                                             TYPE: RNA
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Publication No. US20050153916A1

GENERAL INPORMATION:

APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Beigelman, Leonid

TITLE OF INVENTION: RNA INTERference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA INTERFERENCE: 400/209 (MBHB02-708-C)

FILE REFERENCE: 400/209 (MBHB02-708-C)

CURRENT PILING DATE: 2004-08-20

PRIOR APPLICATION NUMBER: PCT/US03/0408

PRIOR APPLICATION NUMBER: US 60/396,600

PRIOR APPLICATION NUMBER: US 60/396,600

PRIOR PILING DATE: 2002-07-17

PRIOR PLING DATE: 2004-05-24

PRIOR PPLICATION NUMBER: US 10/826,966

PRIOR PPLING DATE: 2004-01-17

PRIOR PPLING DATE: 2004-01-14

PRIOR PILING DATE: 2004-01-14

PRIOR FILING DATE: 2004-01-14

PRIOR FILING DATE: 2004-01-14

PRIOR FILING DATE: 2004-01-14

PRIOR FILING DATE: 2004-01-14
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NAME/KEY: misc_feature
LOCATION: (20) (20) (20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 100.0%; Pred. No. 1e+02; Matches 19; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
NAME/KEY: misc feature
LOCATION: (21) ... (21)
OTHER INFORMATION: n stands for thymidine
US-10-923-330-S86
                                                                                                                         FEATURE:
NAME/KEY: misc_feature
LOCATION: (7)..(7)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                          FEATURE:
NAME/KEY: misc_feature
LOCATION: (11) . (11)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
NAME/KEY: misc_feature
LOCATION: (15)..(19)
OTHER INFORMATION: 2'-deoxy
                                                                                                                                                                                                                                          LOCATION: (8). (10)
OTHER INFORMATION: 2'-deoxy
                                                                             LOCATION: (3). . (6)
OTHER INFORMATION: 2'-deoxy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LOCATION: (12)...(13)
OTHER INFORMATION: 2'-deoxy
                            FEATURE:
NAME/KEY: misc_feature
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; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (21)...(21)
. CTHER INFORMATION: n stands for thymidine
US-10-923-330-588
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LOCATION: (2)...(5)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                 147 TCCACCGTTCATTCTAGAG 165
                                                                                                                                                                                                                                                                                                                                                                                                                                                     19 rccaccerrcarreragae 1
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ORGANISM: Artificial Sequence
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LOCATION: (1). (1)
OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
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NAME/KEY: misc feature
LOCATION: (6). (9)
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Reiselman, Leonic
FILLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400/209 (WHBH02-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT PILLNG DATE: 2004-05-24
FRIOR FILLNG DATE: 2002-07-17
FRIOR PELING DATE: 2002-07-17
FRIOR APPLICATION NUMBER: US 60/396,600
FRIOR APPLICATION NUMBER: US 10/362,966
FRIOR APPLICATION NUMBER: US 10/757,803
FRIOR FILLNG DATE: 2004-04-16
FRIOR PLING DATE: 2004-04-16
FRIOR PLING DATE: 2004-01-14
FRIOR PLING DATE: 2003-11-24
FRIOR FILLNG DATE: 2003-11-24
FRIOR PLING DATE: 2003-11-23
FRIOR PLING DATE: 2003-11-23
FRIOR PLING DATE: 2003-11-23
FRIOR PLING DATE: 2003-05-23
FRIOR APPLICATION NUMBER: US 10/444,853
FRIOR PLING DATE: 2003-05-23
FRIOR PLING DATE: 2003-05-23
FRIOR APPLICATION NUMBER: PCT/US03/05028
FRIOR PLING DATE: 2003-02-20
FRIOR PLING DATE: 2003-02-21
FRIOR PLING DATE: 2003-02-21
FRIOR PLING DATE: 2003-02-20
FRIOR PLING DATE: 2003-02-21
FRIOR PLING DATE: 2003-02-21
FRIOR PLING DATE: 2003-02-23
FRIOR PLING DATE: 2003-02-20
FRIOR PLING DATE: 2003-02-21
FRIOR PLING DATE: 2003-02-21
FRIOR PLING DATE: 2003-02-23
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LOCATION: (20)...(20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
NAME/KEY: misc_feature
LOCATION: (9)..(9)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
NAME/KEY: misc feature
LOCATION: (1). . . (4)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY: misc_feature
LOCATION: (13)..(13)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (16)...(16)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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ORGANISM: Artificial Sequence
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LOCATION: (5). (8)
OTHER INFORMATION: 2'-deoxy
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LOCATION: (10)...(12)
OTHER INFORMATION: 2'-deoxy
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LOCATION: (14)..(15)
OTHER INFORMATION: 2'-deoxy
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STHER INFORMATION: 2'-deoxy
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NAME/KEY: misc_feature
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OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
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 Length 21;
Query Match
4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches 0; Indels
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NAME/KEY: misc feature
LOCATION: (10) (10)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OCHER INFORMATION: n stands for thymidine US-10-923-330-590
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NAME/KEX: misc feature
LOCATION: (15) (15)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (18)...(18)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                           NAME/KEY: misc_feature
LOCATION: (1)._(1)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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Sequence 591, Application US/10923330
Publication No. US20050153916A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              149 CACCGTTCATTCTAGAGCA 167
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NUMBER OF SEQ ID NOS: 768
SOFWARE: Patentin version 3.3
SEQ ID NO 590
LENGTH: 21
                                                                                                                            ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                               NAME/KEY: misc_feature
LOCATION: (2)._(2)
OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: misc_feature
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LOCATION: (11)..(11)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              LOCATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
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US-10-231-330-590/C

US-10-231-330-590/C

Publication No. US20050153916A1

GENERAL INFORMATION:

APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: McSwiggen, James

APPLICANT: McSwiggen, James

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RNA INTERFERENCE: 2004-08-20

CURRENT FILING DATE: 2004-08-20

PRIOR FILING DATE: 2004-09-24

PRIOR FILING DATE: 2004-05-24

PRIOR PILING DATE: 2004-05-24

PRIOR PILING DATE: 2004-01-14

PRIOR PILING DATE: 2004-01-14

PRIOR PILING DATE: 2004-01-14

PRIOR PILING DATE: 2003-11-23

PRIOR PILING DATE: 2003-11-23

PRIOR PILING DATE: 2003-11-23

PRIOR PILING DATE: 2003-11-23

PRIOR PILING DATE: 2003-02-23

PRIOR PILING DATE: 2003-02-20

PRIOR PILING DATE: 2003-02-20
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NAME/KRX: misc feature
LOCATION: (20)-.(20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 100.0%; Pred. No. 1e+02; Matches 19; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: misc_feature

// LOCATION: (21)...(21)

// OTHER INFORMATION: n stands for thymidine

US-10-923-330-589
                                                                                                    FEATURE:
NAME/KEX: misc_feature
LOCATION: (14)-.(14)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                            NAME/KEY: misc feature
LOCATION: (17) ... (17)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          148 CCACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19 CCACCGTTCATTCTAGAGC 1
                                                          LOCATION: (11)..(13)
OTHER INFORMATION: 2'-deoxy
                                                                                                                                                                                                                           NAME/KEY: misc_feature
LOCATION: (15)...(16)
OTHER INFORMATION: 2'-deoxy
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                           LOCATION: (18)...(19)
OTHER INFORMATION: 2'-deoxy
FEATURE:
NAME/KEY: misc feature
LOCATION: (11)...(13)
                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
NAME/KEY: misc_feature
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OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
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OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 100.0%; Pred. No. 1e+02; Matches 19; Conservative 0; Mismatches 0; Indels
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150 ACCGTTCATTCTAGAGCAA 168
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: misc feature
LOCATION: (2)...(3)
OTHER INFORMATION: 2'-deoxy
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LENGTH: 21
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APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-06-20
PRIOR APPLICATION NUMBER: PCT/US03/0408
PRIOR APPLICATION NUMBER: PCT/US03/16390
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-46
PRIOR PRIOR APPLICATION NUMBER: US 10/826,966
PRIOR PRIOR APPLICATION NUMBER: US 10/720,448
PRIOR PILING DATE: 2004-01-14
PRIOR PILING DATE: 2003-11-24
PRIOR APPLICATION NUMBER: US 10/720,448
PRIOR PILING DATE: 2003-11-24
PRIOR APPLICATION NUMBER: US 10/59,059
PRIOR PILING DATE: 2003-11-24
PRIOR APPLICATION NUMBER: US 10/693,059
PRIOR PILING DATE: 2003-01-23
PRIOR APPLICATION NUMBER: PCT/US03/05346
PRIOR PILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR PILING DATE: 2003-02-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: misc feature
LOCATION: (1). (2)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (4)._(7)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (12)...(12)
JTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (16)...(16)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (19)..(19)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
FEATURE:
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ORGANISM: Artificial Sequence
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LOCATION: (3)...(3)
OTHER INFORMATION: 2'-deoxy
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LOCATION: (13)...(15)
OTHER INFORMATION: 2'-deoxy
FEATURE:
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LOCATION: (8)._(11)
OTHER INFORMATION: 2'-deoxy
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LOCATION: (17)...(18)
OTHER INFORMATION: 2'-deoxy
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LENGTH: 21
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9 Sequence 502, Application US/10923330
9 Publication No. US20050153916A1
GENERAL INFORMATION:
19 APPLICANT: Sitna Therapeutics, Inc.
19 APPLICANT: McSusgen, James
19 APPLICANT: McSusgen, James
19 APPLICANT: McSusgen, James
10 APPLICANT: MVBER: US/10/923,330
10 FILE REFRENCE: 400/209 (MSHB2-708-C)
10 FILE REFRENCE: 400/209 (MSHB2-708-C)
10 FILE REPRICATION NUMBER: PCT/US04/16390
10 FRIOR FILING DATE: 2002-07-17
10 FRIOR FILING DATE: 2002-07-17
10 FRIOR FILING DATE: 2003-07-17
11 FRIOR FILING DATE: 2003-07-17
12 FRIOR FILING DATE: 2003-01-14
13 FRIOR FILING DATE: 2003-11-24
14 FRIOR FILING DATE: 2003-11-24
15 FRIOR FILING DATE: 2003-11-24
16 FRIOR FILING DATE: 2003-11-24
16 FRIOR FILING DATE: 2003-11-23
16 FRIOR FILING DATE: 2003-11-23
17 FRIOR FILING DATE: 2003-11-23
18 FRIOR FILING DATE: 2003-05-24
18 FRIOR FILING DATE: 2003-05-23
18 FRIOR FILING DATE: 2003-05-23
18 FRIOR FILING DATE: 2003-05-24
18 FRIOR FILING DATE: 2003-05-23
18 FRIOR FILING DATE: 2003-05-23
18 FRIOR FILING DATE: 2003-05-34
18 FRIOR FILING DATE: 2003-05-23
18 FRIOR FILING DATE: 2003-05-34
18 FRIOR FILING DATE: 2003-05-35
18 FRIOR
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NAME/KEY: misc_feature
LOCATION: (1)..(1)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY: misc_feature
LOCATION: (20)...(20)
OTHER INPORMATION: Phosphorothioate 3'-Internucleotide Linkage
FEATURE:
NAME/KEY: misc_feature
LOCATION: (21)...(21)
OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 100.0%; Pred. No. 1e+02; Matches 19; Conservative 0; Mismatches 0; Indels
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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ORGANISM: Artificial Sequence
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JOSEPHONE STATES STATES SATIONARY SEQUENCE 594, Application US/10923330

JOSEPHONE STATE Therapeutics, Inc.

JAPPLICANT: SLIVE THE Therapeutics, Inc.

JAPPLICANT: More Application Sequence States Stat
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LOCATION: (20) ...(20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
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NAME/KEY: misc_feature
LOCATION: (21)...(21)
US.-10-923-330-592
                                                                                                                                 NAME/KEY: misc feature
LOCATION: (9). (11)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY: misc_feature
LOCATION: (12)...(13)
OTHER INFORMATION: 2'-deoxy
FEATURE:
NAME/KEY: misc_feature
LOCATION: (5)._(8)
OTHER INFORMATION: 2'-deoxy
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LENGTH: 21
TYPE: RNA
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Publication No. US20050153916A1
GENERAL INCOMPATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-08-20
PRIOR FILING DATE: 2004-05-24
  siNA sense region
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                                           NAME/KEY: misc_feature
LOCATION: (1)._(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
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CTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity US-10-923-330-594
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 63.2%; Pred. No. 1e+02; Matches 12; Conservative 7; Mismatches 0; Indels
OTHER INFORMATION: Description of Artificial Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LOCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                 LOCATION: (7). (8)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LOCATION: (10) ... (12)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                LOCATION: (1). (5)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (9)._(9)
OTHER INFORMATION: 2'-O-methyl
                                                                                                                                                                                                                                                            LOCATION: (6). (6)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LOCATION: (18)...(19)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LOCATION: (13)...(13)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
NAME/KEY: misc_feature
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NAME/KEY: misc_feature
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NAME/KEY: misc_feature
                                                                                                                                     NAME/KEY: misc_feature
                                                                                                                                                                                                                                 NAME/KEY: misc feature
LOCATION: (6)..(6)
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LOCATION: (14)..(17)
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OTHER INFORMATION: Description of Artificial Sequence: siNA sense region
                                    PRIOR PLING DATE: 2002-07-17

PRIOR PLING DATE: 2002-07-17

PRIOR PLING DATE: 2002-07-17

PRIOR PLING DATE: 2004-05-24

PRIOR PLING DATE: 2004-04-16

PRIOR APPLICATION NUMBER: US 10/757,803

PRIOR PLING DATE: 2004-04-16

PRIOR PLING DATE: 2004-01-124

PRIOR PLING DATE: 2003-11-24

PRIOR PLING DATE: 2003-11-24

PRIOR PLING DATE: 2003-11-23

PRIOR PLING DATE: 2003-11-23

PRIOR PPLICATION NUMBER: US 10/444,853

PRIOR PPLICATION NUMBER: PCT/US03/05946

PRIOR PLING DATE: 2003-05-220

PRIOR PLING DATE: 2003-02-20

PRIOR PLING DATE: 2003-02-20

PRIOR PLING DATE: 2003-02-20

PRIOR APPLICATION NUMBER: PCT/US03/05028

PRIOR PLING DATE: 2003-02-20

PRIOR PLING DATE: 2003-02-20

PRIOR APPLICATION NUMBER: PCT/US03/05028

PRIOR APPLICATION NUMBER: PCT/US03/05028
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NAME:
NAM
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NAME/KEY: misc feature
LOCATION: (21)...(21)
OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-595
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: misc feature
LOCATION: (20) ... (20)
OTHER INFORMATION: n stands for thymidine
APPLICATION NUMBER: US 60/396,600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: misc_feature
LOCATION: (1)._(4)
OTHER_INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (6)._(7)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (13)...(16)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-0-methyl
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OTHER INFORMATION: 2'-0-methyl
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: misc feature
LOCATION: (17)...(19)
OTHER INFORMATION: 2'-O-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: misc_feature
LOCATION: (12)...(12)
OTHER INFORMATION: 2'-0-methyl
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LOCATION: (8)..(8)
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RESULT 1310-596

Sequence 586, Application US/10923330

Sequence 586, Application US/10923330

Sequence 586, Application US/10923330

GENERAL INFORMATION:

APPLICANT: SITE AT The Tapecutics, Inc.

APPLICANT: Begglann, Jeneid

TITLE OF INVENTION: EARL Leonid

TITLE OF INVENTION: EARL LEONID

TITLE OF INVENTION: EARL OF 1004-08-20

CURRENT APPLICATION NUMBER: DC10/10/923, 30

FILE REFERENCE: 400/209 (MBHB02-109-C)

CURRENT APPLICATION NUMBER: DC10/10/923, 30

PRIOR APPLICATION NUMBER: DC10/10/920, 330

PRIOR APPLICATION NUMBER: DC10/10/920, 600

PRIOR PELING DATE: 2004-05-24

SPRIOR APPLICATION NUMBER: US 10/826, 966

PRIOR PELING DATE: 2004-01-14

PRIOR PELING DATE: 2004-01-14

PRIOR PELING DATE: 2004-01-14

PRIOR PELING DATE: 2004-01-14

PRIOR PELING DATE: 2003-11-23

PRIOR PELING DATE: 2003-11-23

PRIOR APPLICATION NUMBER: US 10/693, 059

PRIOR APPLICATION NUMBER: PCT/US03/05346

PRIOR APPLICATION NUMBER: PCT/US03/05346

PRIOR APPLICATION NUMBER: PCT/US03/05346

PRIOR APPLICATION NUMBER: PCT/US03/05346

PRIOR PRIOR PRIOR PAPE: 2003-00-20

PRIOR PRIOR PRIOR PAPE: 2003-00-20

PRIOR PRIOR PRIOR PAPE: 2003-00-20

PRIOR PAPE: 2003-0
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                                                                                                        Gaps
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OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
NAME/KEY: misc_feature
LOCATION: (1)...(3)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 63.2%; Pred. No. 1e+02; Matches 12; Conservative 7; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: misc feature
LOCATION: (5).7(6)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                        146 TICCACCGTICATICIAGA 164
                                                                                                                                                                                                            1 UUCCACCGUUCAUUCUAGA 19
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OTHER INFORMATION: 2'-0-methyl
FEATURE:
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LOCATION: (7)..(7)
OTHER INFORMATION: 2'-O-methyl
FEATURE:
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
NAME/KEY: misc_feature
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NAME/KEY: misc_feature
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APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400/209 (WBHB02-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
                                  sina sense region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
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                                                                                                                           LOCATION: (1). (1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 21;
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FEATURE:
OTHER INFORMATION: Description of Artificial Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ouery Match
4.2%; Score 19; DB 1;
Best Local Similarity 73.7%; Pred. No. 1e+02;
Matches 14; Conservative 5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LOCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LOCATION: (4). (5)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LOCATION: (11)...(14)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                LOCATION: (1)._(2)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           148 CCACCGTTCATTCTAGAGC 166
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: misc feature
LOCATION: (10)...(10)
OTHER INFORMATION: 2'-O-methyl
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OTHER INFORMATION: 2'-O-methyl
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OTHER INFORMATION: 2'-0-methyl
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OTHER INFORMATION: 2'-O-methyl
                                                                                           NAME/KEY: misc_feature
                                                                                                                                                                                                                NAME/KEY: misc_feature
LOCATION: (1)...(2)
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LOCATION: (6).7(6)
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LOCATION: (7)...(9)
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USA-10-293-390-597

USA-10-293-390-597

Publication No. U82005015391641

Sequence 597, Application US/10923330

Sepulation No. U82005015391641

APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: Garway Garm, James

APPLICANT: MANGARGEN, Leonid

ITILE OF INVENTION: RAPECESTON USING Short Interfering RNA (siNA)

FILE REPERRECE: 400/209 (MHB02-708-C)

FURRENT FILING DATE: 2004-08-24

PRIOR PELLOR DATE: 2004-06-24

PRIOR PELLOR DATE: 2002-07-17

PRIOR PELLOR DATE: 2002-07-17

PRIOR PELLOR DATE: 2002-07-17

PRIOR PELLOR WIMBER: US 60/396,600

PRIOR FILING DATE: 2002-04-16

PRIOR PELLOR TON NUMBER: US 10/757,803

PRIOR PELLOR TON NUMBER: US 10/757,803

PRIOR PELLOR DATE: 2003-11-24

PRIOR PELLOR DATE: 2003-02-20

PRIOR APPLICATION NUMBER: US 10/693,059

PRIOR APPLICATION NUMBER: PET/US03/05028

PRIOR PELLOR TON NUMBER: PET/US03/05028

PRIOR APPLICATION NUMBER: PET/US03/05028

PRIOR PELLOR TON NUMBER: PET/US03/05028

PRIOR PELLOR NUMBER: PET/US03/05028

PRIOR PELLOR NUMBER: PET/US03/05028

PRIOR PELLOR PET/US03/05028

PRIOR PELLOR NUMBER: PET/US03/05028

PROPER PET/US03/05028

PET/US03/05028

PET/US03/05028

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PET/US03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: misc_feature

| LOCATION: (21)...(21)

| OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity

US-10-923-330-596
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 19; DB 1; Length 21;
Pred. No. 1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LOCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                FEATURE:
NAME/KEY: misc_feature
LOCATION: (12) ... (15)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
   LOCATION: (8)..(10)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  147 TCCACCGTTCATTCTAGAG 165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
NAME/KEY: misc_feature
LOCATION: (16)..(19)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                           LOCATION: (11)...(11)
OTHER INFORMATION: 2'-O-methyl
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 68.4
Matches 13; Conservative
                                                                                           NAME/KEY: misc feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: misc_feature
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Gaps

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OTHER INFORMATION: Description of Artificial Sequence: sink sense region
PRIOR APPLICATION NUMBER: PCT/US03/04088

PRIOR PILING DATE: 2004-08-20

PRIOR FILING DATE: 2004-05-24

PRIOR PELLON NUMBER: US 60/396,600

PRIOR PELLON NUMBER: US 60/396,600

PRIOR PELLON NUMBER: US 10/826,966

PRIOR APPLICATION NUMBER: US 10/826,966

PRIOR APPLICATION NUMBER: US 10/757,803

PRIOR FILING DATE: 2004-01-14

PRIOR FILING DATE: 2004-01-14

PRIOR FILING DATE: 2004-01-14

PRIOR FILING DATE: 2003-11-24

PRIOR PELLON NUMBER: US 10/693,059

PRIOR FILING DATE: 2003-11-24

PRIOR PILING DATE: 2003-11-24

PRIOR PILING DATE: 2003-05-23

PRIOR FILING DATE: 2003-05-23

PRIOR FILING DATE: 2003-05-23

PRIOR FILING DATE: 2003-05-23

PRIOR FILING DATE: 2003-05-23

PRIOR PRIOR APPLICATION NUMBER: PCT/US03/05346

PRIOR PRIOR APPLICATION NUMBER: PCT/US03/05346

PRIOR PRIOR PRIOR DATE: 2003-05-23

PRIOR FILING DATE: 2003-05-23

PRIOR PRIOR PRIOR PRIOR NUMBER: PCT/US03/0508

PRIOR PRI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: misc feature
LOCATION: (1)...(1)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: misc feature
LOCATION: (3). (4)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
NAME/KEY: misc_feature
LOCATION: (18)..(18)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY: misc_feature
LOCATION: (9)..(9)
OTHER INFORMATION: 2'-O-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: misc_feature
LOCATION: (2)...(2)
OTHER INFORMATION: 2'-0-methyl
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LOCATION: (5)._(5)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: misc feature
LOCATION: (14)...(17)
OTHER INFORMATION: 2'-O-methyl
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LOCATION: (19)..(19)
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Sequence 599, Application US/10923330
| Sequence 599, Application US/10923330
| Publication No. US205050153916A1
| GENERAL INFORMATION:
| APPLICANT: SITMA THERAPRUATION:
| APPLICANT: MCSWAGGEN, James
| APPLICANTON WUMBER: US/10/923, 330
| CURRENT PAPLICATION WUMBER: US/10/923, 330
| CURRENT PAPLICATION WUMBER: US/10/923, 330
| CURRENT APPLICATION WUMBER: US/10/926, 966
| PRIOR APPLICATION WUMBER: US/10/926, 966
| PRIOR APPLICATION WUMBER: US/10/720, 448
| PRIOR PELICATION WUMBER: US/10/93, 059
| PRIOR PELICATION WUMBER: US/10/93,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .
                                                                    NAME/KEY: misc feature
LOCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
FEATURE:
NAME/KEY: misc feature
LOCATION: (21)...(21)
OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LOCATION: (1)..(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
                                                                                                                                                                                                                                                                                                                                                                                                                             4.2%; Score 19; DB 1; Length 21; 73.7%; Pred. No. 1e+02; tive 5; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  149 CACCGTTCATTCTAGAGCA 167
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OTHER INFORMATION: 2'-0-methyl
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LOCATION: (1)._(1)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 73.7
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: misc_feature
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NAME/KEY: misc_feature
LOCATION: (2)...(3)
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siNA sense region
                PRIOR FILING DATE: 2003-11-23
PRIOR FILING DATE: 2003-10-23
PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR FILING DATE: 2003-05-23
PRIOR FILING DATE: 2003-02-20
PRIOR PILING DATE: 2003-02-20
PRIOR APPLICATION WUMBER: PCT/US03/0508
PRIOR APPLICATION WUMBER: PCT/US03/05028
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 768
SOFTWARE: Patentin version 3.3
SEQ ID NO 600
LENGTH: 21
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| LOCATION: (21)...(21)

| CTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity

US-10-923-330-600
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LOCATION: (1). 7(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 21;
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                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Description of Artificial Sequence:
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4.2%; Score 19; DB 1;
Best Local Similarity 73.7%; Pred. No. 1e+02;
Matches 14; Conservative 5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: misc feature
LOCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
APPLICATION NUMBER: US 10/693,059
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LOCATION: (17)..(18)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LOCATION: (7). (8)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GAAGAGUUGGGCUCUGUCA 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LOCATION: (1). (6)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LOCATION: (9). (11)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LOCATION: (16)...(16)
OTHER INFORMATION: 2'-0-methy]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LOCATION: (19)...(19)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                   TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc feature
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NAME/KEY: misc_feature
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US-10-923-330-602/c
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publication No. US20050153916A1

Sequence 600, Application US/10923330

sublication No. US20050153916A1

GENERAL INFORMATION:

APPLICANT: Siran Therapeutics, Inc.

APPLICANT: Beigelman, Leonid

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RNA INTERFERENCE: 400/209 (MBHB02-708-C)

FILE REFERENCE: 400/209 (MBHB02-708-C)

FURRENT APPLICATION NUMBER: US/10/923,330

CURRENT APPLICATION NUMBER: DCT/US03/0408

PRIOR FILING DATE: 2004-05-24

PRIOR PLING DATE: 2004-05-24

PRIOR PLING DATE: 2004-06-24

PRIOR FILING DATE: 2004-04-16

PRIOR FILING DATE: 2004-04-16

PRIOR FILING DATE: 2004-01-14

PRIOR FILING DATE: 2004-01-14
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; LOCATION: (21)...(21)
; OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-599
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 73.7%; Pred. No. 1e+02; Matches 14; Conservative 5; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
NAME/KEY: misc_feature
LOCATION: (20)..(20)
OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                 FEATURE:
NAME/KEY: misc_feature
LOCATION: (9)..(12)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                             NAME/KEY: misc_feature
LOCATION: (5). (7)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
NAME/KEY: misc_feature
LOCATION: (17)..(17)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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                                                NAME/KEY: misc_feature
LOCATION: (4)._(4)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc_feature
LOCATION: (18)...(19)
OTHER INFORMATION: 2'-O-methyl
                                                                                                                                                                                                                                                      NAME/KEY: misc_feature
LOCATION: (8).7(8)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LOCATION: (13)..(16)
OTHER INFORMATION: 2'-0-methyl
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FEATURE:
NAME/KEY: misc_feature
LOCATION: (1) ...(3)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                    145 CTTCCACCGTTCATTCTAG 163
                                                                                                                                                                                                                                                                                                                                                                                                      19 CTTCCACCGTTCATTCTAG 1
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LOCATION: (4). (7)
OTHER INFORMATION: 2'-0-methyl
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ORGANISM: Artificial Sequence
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APPLICANT: SITUR TIRETAPEULICS, INC.
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leones
APPLICANT: Beigelman, Leones
APPLICANT: Beigelman, Leones
TITLE OF INVENTION: RAY Interference Mediated Inhibition of Telomerase Gene
FILE REFERENCE: 400/209 (MRHB02-108-C)
CURRENT APPLICATION NUMBER: 108/10/923,330
CURRENT FILING DATE: 2004-08-20
PRIOR APPLICATION NUMBER: PCT/US03/0408
PRIOR APPLICATION NUMBER: PCT/US03/0408
PRIOR APPLICATION NUMBER: PCT/US04/16390
PRIOR PELING DATE: 2004-05-24
PRIOR PELING DATE: 2004-01-14
PRIOR PELING DATE: 2004-01-14
PRIOR PELING DATE: 2004-01-14
PRIOR PELING DATE: 2003-01-14
PRIOR PELING DATE: 2003-11-24
PRIOR PELING DATE: 2003-05-23
PRIOR PELING DATE: 2003-05-20
PRIOR PELING DATE: 2003-05-30
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LOCATION: (20)...(20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: misc_feature
LOCATION: (1)._(2)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (11) ... (11)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (14)...(14)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-0-methyl
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NAME/KEY: misc_feature
LOCATION: (8)..(10)
OTHER_INFORMATION: 2'-0-methyl
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NAME/KEY: misc_feature
LOCATION: (12)..(13)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: RNA
ORGANISM: Artificial Sequence
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INFORMATION: 2'-O-methyl
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LOCATION: (7)...(7)
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LENGTH: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OCATION:
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Sequence 603, Application US/1092330
| Publication NO. US20050153916A1
| Publication NO. US20050153916A1
| Publication NO. US20050153916A1
| APPLICANT: Gravalgen, James
| APPLICANT: Beigelman, Leonid
| TITLE OF INVENTION: RAN Interference Mediated Inhibition of Telomerase Gene
| TITLE OF INVENTION: RAN Interference Mediated Inhibition of Telomerase Gene
| TITLE OF INVENTION: RAN Interference Mediated Inhibition of Telomerase Gene
| TITLE OF INVENTION: RAN INTERFERENCE: 400/209 (MRHE02-708-C)
| CURRENT FELLING DATE: 2004-08-24
| FRIOR PAPLICATION NUMBER: DS/10/923,330
| CURRENT FILING DATE: 2004-05-24
| PRIOR PAPLICATION NUMBER: DS/10/926,600
| PRIOR FILING DATE: 2004-05-24
| PRIOR PAPLICATION NUMBER: US 10/750,966
| PRIOR PAPLICATION NUMBER: US 10/750,969
| PRIOR PAPLICATION NUMBER: US 10/750,969
| PRIOR PAPLICATION NUMBER: US 10/720,446
| PRIOR PAPLICATION NUMBER: US 10/720,969
| PRIOR PAPLICATION NUMBER: PET/US03/05026
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
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                                                                                                                                                                                                                Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 100.0%; Pred. No. 16+02; Matches 19; Conservative 0; Mismatches 0; Indels
; NAME/KEY: misc_feature
; LOCATION: (21)...(21)
; OTHER INFORMATION: n stands for thymidine
US-10-923-330-602
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: misc feature
LOCATION: (9)...(11)
OTHER INFORMATION: 2'-O-methyl
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OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Baigalman, Leonid
TITLE DE INVENTION: Baigalman, Leonid
TITLE OF INVENTION: BAY Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: BAY Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: BAY Interference Mediated Interfering RNA (siNA)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-08-20
PRIOR PILING DATE: 2004-08-24
PRIOR APPLICATION NUMBER: US 60/396,600
PRIOR FILING DATE: 2002-07-17
PRIOR APPLICATION NUMBER: PCT/US04/16390
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NAME/KEY: misc_feature
LOCATION: (20) ... (20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 100.0%; Pred. No. 1e+02; Matches 19; Conservative 0; Mismatches 0; Indels
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NAME/KEY: misc feature
LOCATION: (21)...(21)
OTHER INFORMATION: n stands for thymidine
US-10-923-330-604
                                                                                                                                                                                                                                                                                                                                                                FEATURE:
NAME/KRY: misc_feature
LOCATION: (9)..(9)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                 LOCATION: (1). (4)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LOCATION: (10)...(12)
OTHER INFORMATION: 2'-O-methyl
FEATURE:
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OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LOCATION: (17)..(19)
OTHER INFORMATION: 2'-0-methyl
   ORGANISM: Artificial Sequence
                                                                                                                                                                                                                              FEATURE:
NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
NAME/KEY: misc_feature
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LOCATION: (14)..(15)
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LOCATION: (16)..(16)
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LOCATION: (17)..(19)
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GENERAL INCRAMINES.

GENERAL INCRAMINES.

APPLICANT: SITUR Therapeutice, Inc.

APPLICANT: McSa'ggan, James

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

FILE REFRENCE: 400/209 (MHHBOL-708-C)

CURRENT PILING DATE: 2004-08-20

FRIOR PILING DATE: 2004-08-20

FRIOR PILING DATE: 2004-05-34

FRIOR FILING DATE: 2004-05-34

FRIOR FILING DATE: 2004-06-24

FRIOR FILING DATE: 2004-04-16

FRIOR FILING DATE: 2004-04-16

FRIOR FILING DATE: 2004-04-16

FRIOR FILING DATE: 2004-04-16

FRIOR FILING DATE: 2003-11-34

FRIOR FILING DATE: 2003-05-33

FRIOR FILING DATE: 2003-05-33

FRIOR FILING DATE: 2003-05-30

FRIOR APPLICATION NUMBER: FCT/US03/05028

FRIOR FILING DATE: 2003-05-30

FRIOR FILI
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LOCATION: (20) ... (20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1; Length 21;
. 1e+02;
ches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 4.2%; Score 19; DB Best Local Similarity 100.0%; Pred. No. 1e+ Matches 19; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; LOCATION: (21)...(21); OTHER INFORMATION: n stands for thymidine US-10-923-330-603
   NAME/KEY: misc_feature
LOCATION: (12) .. (12)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                              FEATURE:
NAME/KEX: misc_feature
LOCATION: (15)-.(15)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 604, Application US/10923330; Publication No. US20050153916A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    146 TTCCACCGTTCATTCTAGA 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TICCACCGTICATICIAGA 1
                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: misc_feature
LOCATION: (16)..(19)
OTHER INFORMATION: 2'-O-methyl
                                                                                                   FEATURE:
NAME/KEY: misc_feature
LOCATION: (13)..(14)
OTHER_INFORMATION: 2'-O-methyl
misc_feature
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US-10-923-330-604/c
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Сарв

Page 82

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Sequence 606, Application US/10923330
| Sequence 606, Application US/10923330
| Publication No. US20050153916A1
| GENERAL INFORMATION.
| PAPLICANT: Sirma Therapeutics, Inc.
| APPLICANT: McSwigen, James
| FRICK FILING DATE: 2004-06-24
| FRICK FILING DATE: 2004-06-24
| FRICK FILING DATE: 2004-01-16
| FRICK FILING DATE: 2003-01-12
| FRICK FILING DATE: 2003-02-20
| FRICK FILING DATE: 2003-03-23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY:
LOCATION: (1)._(1)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY: misc_feature
LOCATION: (3)..(6)
OTHER_INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
NAME/KEY: misc_feature
LOCATION: (11)...(11)
GTHER INFORMATION: 2'-deoxy-2'-fluoro
FEATURE:
                                                                       148 CCACCGTTCATTCTAGAGC 166
                                                                                                                     19 CCACCGTTCATTCTAGAGC 1
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NAME/KEY: misc_feature
LOCATION: (7)..(10)
OTHER INFORMATION: 2'-0-methyl
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NAME/KEY: misc_feature
LOCATION: (2)...(2)
OTHER INFORMATION: 2'-0-methyl
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LOCATION: (12)...(14)
OTHER INFORMATION: 2'-0-methyl
FEATURE:
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                               US-10-923-330-606/c
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OTHER INFORMATION: Description of Artificial Sequence: siNA antisense region
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PRIOR FILING DATE: 2004-05-24

PRIOR PILING DATE: 2004-01-16

PRIOR APPLICATION NUMBER: US 10/826,966

PRIOR PILING DATE: 2004-01-16

PRIOR PILING DATE: 2004-01-14

PRIOR PILING DATE: 2004-01-14

PRIOR PILING DATE: 2003-11-23

PRIOR FILING DATE: 2003-11-23

PRIOR PILING DATE: 2003-11-23

PRIOR PILING DATE: 2003-11-23

PRIOR APPLICATION NUMBER: US 10/44,853

PRIOR PILING DATE: 2003-05-23

PRIOR PILING DATE: 2003-02-20

SOFTWARE: PALENT APPLICATION NUMBER: PCT/US03/05028

PRIOR PILING DATE: 2003-02-20

SOFTWARE: PALENT APPLICATION NUMBER: PCT/US03/05028

PRIOR PILING DATE: 2003-02-20

PRIOR PILING DATE: 2003-02-20
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LOCATION: (20) (20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:

NAME/KEY: misc_feature

COTATION: (21)...(21)

OTHER INFORMATION: n stands for thymidine
US-10-923-330-605
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(1)
OTHER INFORMATION: 2'-O-methyl
FEATURE:
NAME/KEY: misc_feature
LOCATION: (2)...(5)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY: misc_feature
LOCATION: (14)..(14)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (17)...(17)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
NAME/KEY: misc_feature
LOCATION: (6)..(9)
OTHER INFORMATION: 2'-O-methyl
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NAME/KEY: misc_feature
LOCATION: (11)...(13)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: misc feature
LOCATION: (18)...(19)
OTHER INFORMATION: 2'-0-methyl
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OTHER INFORMATION: 2'-0-methyl
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LOCATION: (15)..(16)
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OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
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Publication No. US20050153916A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Beigelman, Leonid
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-08-20
PRIOR FILING DATE: 2004-08-24
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OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
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4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
NAME/KEX: misc_feature
LOCATION: (19)..(19)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                         LOCATION: (4)...(/)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LOCATION: (12)...(12)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                           LOCATION: (1)._(2)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: 2'-deoxy-2'-fluoro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: misc_feature
LOCATION: (17)..(18)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc feature
LOCATION: (8). (11)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LOCATION: (13). (15)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: 2'-0-methyl
ORGANISM: Artificial Sequence
                                                                                                                             NAME/KEY: misc_feature
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NAME/KEY: misc_feature
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LOCATION: (16)..(16)
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                                                                                                                                                                                                                                                                                               COCATION:
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LOCATION: (20) ... (20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
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Pred. No. 1e+02;
0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 100.0%; Pred. No. 1e+
Matches 19; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc_feature

// LOCATION: (21)...(21)

OTHER INFORMATION: n stands for thymidine

US-10-923-330-606
NAME/KEY: misc feature
LOCATION: (15)...(15)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                        FEATURE:
NAME/KEY: misc_feature
LOCATION: (18)..(18)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 607, Application US/10923330; Publication No. US20050153916A1; GENERAL INFORMATION:
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                                                                                                                             NAME/KEY: misc feature
LOCATION: (16)...(17)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: misc_feature
LOCATION: (19)...(19)
OTHER INFORMATION: 2'-0-methyl
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FEATURE:
NAME/KEY: misc_feature
LOCATION: (20)
OTHER INFORMATION: n stands for thymidine
                                                               US-10-923-330-610
; Sequence 610, Application US/10923330
; Publication No. US20050153916A1
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ORGANISM: Artificial Sequence
FEATURE:
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                  PRIOR FILING DATE: 2002-07-17

PRIOR FILING DATE: 2002-07-17

PRIOR APPLICATION NUMBER: PCT/US04/16390

PRIOR PILING DATE: 2004-05-24

PRIOR APPLICATION NUMBER: US 10/826,966

PRIOR PELING DATE: 2004-04-16

PRIOR PELING DATE: 2004-01-12

PRIOR PELING DATE: 2004-01-124

PRIOR PELING DATE: 2003-11-24

PRIOR PELING DATE: 2003-11-24

PRIOR APPLICATION NUMBER: US 10/693,059

PRIOR PELING DATE: 2003-11-24

PRIOR PELING DATE: 2003-05-23

PRIOR PELING DATE: 2003-05-23

PRIOR PELING DATE: 2003-05-23

PRIOR PELING DATE: 2003-05-23

PRIOR APPLICATION NUMBER: PCT/US03/0508

PRIOR APPLICATION NUMBER: PCT/US03/0508

PRIOR APPLICATION NUMBER: PCT/US03/0508

PRIOR APPLICATION NUMBER: PCT/US03/0508

PRIOR PELING DATE: 2003-02-20

PROMBER OF SEQ ID NOS: 768

SOFTWARE: PATENTIN VERSION 3.3

FENDING OF 08
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NAME/KEY: misc_feature
LOCATION: (20)...(20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4.2%; Score 19; DB 1; Length 21; 100.0%; Pred. No. 1e+02; Live 0; Mismatches 0; Indels
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OTHER INFORMATION: n stands for thymidine
APPLICATION NUMBER: US 60/396,600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)._(1)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: misc_feature
LOCATION: (4)...(4)
OTHER_INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: misc_feature
LOCATION: (14)...(19)
OTHER_INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY: misc_feature
LOCATION: (2)...(3)
OTHER INFORMATION: 2'-O-methyl
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LOCATION: (5)._(8)
OTHER INFORMATION: 2'-O-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LOCATION: (12)...(13)
OTHER INFORMATION: 2'-O-methyl
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence
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Best Local Similarity 100.(
Matches 19; Conservative
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NAME/KEY: misc_feature
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NAME/KEY: misc_feature
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GENERAL INFORMATION:

APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REPERRNCE: 400/209 (MBHB02-708-C)
CURRENT FILING DATE: 2004-08-20
FRIOR APPLICATION NUMBER: US 60/396,600
FRIOR PILING DATE: 2002-07-14
FRIOR FILING DATE: 2004-05-24
FRIOR PLING DATE: 2004-06-24
FRIOR PLING DATE: 2004-01-14
FRIOR PLING DATE: 2004-01-14
FRIOR APPLICATION NUMBER: US 10/757,803
FRIOR PLING DATE: 2003-11-23
FRIOR PLING DATE: 2003-11-23
FRIOR PLING DATE: 2003-11-23
FRIOR PLING DATE: 2003-10-20
FRIOR APPLICATION NUMBER: US 10/757,803
FRIOR PLING DATE: 2003-01-23
FRIOR FILING DATE: 2003-02-20
FRIOR APPLICATION NUMBER: US 10/750,488
FRIOR PLING DATE: 2003-02-20
FRIOR APPLICATION NUMBER: US 10/7503/05346
FRIOR FILING DATE: 2003-02-20
FRIOR APPLICATION NUMBER: US CT/US03/05028
FRIOR FILING DATE: 2003-02-20
FRIOR APPLICATION NUMBER: PCT/US03/05028
FRIOR FILING DATE: 2003-02-20
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NAME/KEY: misc_feature
LOCATION: (1)._(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: misc feature; LOCATION: (21)...(21); OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity US-10-923-330-610
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4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 63.2%; Pred. No. 1e+02;
Matches 12; Conservative 7; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 194
US-10-923-330-611
Sequence 611, Application US/10923330
Publication No. US20050153916A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
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APPLICATION NUMBER: US 60/396,600
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          TITLE OF INVENTION: RAA Interference Mediated Inhibition of Telomerase Gene TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA) FILE REFERENCE: 400/200 (Wahlsold-708-C) CURRENT APPLICATION NUMBER: US/10/923,330 CURRENT PAPLICATION NUMBER: PCT/US03/04088 PRIOR PELICATION NUMBER: PCT/US03/04088 PRIOR PELICATION NUMBER: PCT/US03/04088 PRIOR PELICATION NUMBER: US 60/396,600 PRIOR PELICATION NUMBER: US 60/396,600 PRIOR PELICATION NUMBER: US 10/826,966 PRIOR PELICATION NUMBER: US 10/826,966 PRIOR PELICATION NUMBER: US 10/720,448 PRIOR PELICATION NUMBER: US 10/720,448 PRIOR PELICATION NUMBER: US 10/720,448 PRIOR PELICATION NUMBER: US 10/693,059 PRIOR PELICATION NUMBER: PCT/US03/05028 PRIOR PELICATION POLOR: PALM.
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Publication No. US20050153916A1
GENERAL INFORMATION
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400/209 (MBHB02-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-08-20
CURRENT PELICATION NUMBER: PCT/US03/04088
PRIOR FILING DATE: 2004-05-24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PEATURE:
NAME/KEX: misc feature
LOCATION: (21) (21)
OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4.2%; Score 19; DB 1; Length 21; 63.2%; Pred. No. 1e+02;
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LOCATION: (20)..(20)
OTHER INFORMATION: n stands for thymidine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 UUCCACCGUUCAUCUAGA 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: RNA
ORGANISM: Artificial Sequence
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Matches 12, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
NAME/KEY: misc_feature
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US-10-923-330-612
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-10-923-330-611
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LENGTH: 21
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OTHER INFORMATION: Description of Artificial Sequence: siNA sense region
PRIOR FILING DATE: 2002-07-17
PRIOR PLING DATE: 2002-07-17
PRIOR PLING DATE: 2002-07-17
PRIOR PLING DATE: 2004-05-24
PRIOR PLING DATE: 2004-06-24
PRIOR PLING DATE: 2004-06-16
PRIOR PLING DATE: 2004-01-16
PRIOR PLING DATE: 2004-01-14
PRIOR PLING DATE: 2004-01-14
PRIOR PLING DATE: 2003-11-24
PRIOR PLING DATE: 2003-11-23
PRIOR PLING DATE: 2003-11-23
PRIOR PLING DATE: 2003-05-23
PRIOR PLING DATE: 2003-05-23
PRIOR PLING DATE: 2003-02-20
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APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: MedSwiggen, James

APPLICANT: Beigelman, Leonid

TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telo

TITLE OF INVENTION: Expression Using Short Interfering RNA (sin

TITLE OF INVENTION: Expression Using Short Interfering RNA (sin

FILE REFERENCE: 400/209 (MBHB02-708-C)

CURRENT APPLICATION NUMBER: US/10/923,330

PRIOR PELING DATE: 2004-06-24

PRIOR PELING DATE: 2004-06-24

PRIOR APPLICATION NUMBER: US 60/396,600

PRIOR PELING DATE: 2004-06-24

PRIOR FILING DATE: 2004-06-24

PRIOR FILING DATE: 2004-06-24

PRIOR FILING DATE: 2004-04-16

PRIOR FILING DATE: 2004-04-16

PRIOR FILING DATE: 2004-04-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
NAME/KEY: misc_feature
LOMG/KEY: misc_feature
LOMG/TON: (1)..(1)
OTHER INFORMATION: 5.-3 attached terminal deoxyabasic moeity

j LOCATION: (21)...(21)

j OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-612

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LOCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 613, Application US/10923330 Publication No. US20050153916A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       147 TCCACCGTTCATTCTAGAG 165
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: RNA
ORGANISM: Artificial Sequence
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Best Local Similarity 68.41
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
NAME/KEY: misc_feature
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NAME/KEY: misc feature
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PRIOR APPLICATION NUMBER: US 10/720,448
PRIOR PILING DATE: 2003-11-24
PRIOR PELING DATE: 2003-11-24
PRIOR APPLICATION NUMBER: US 10/693,059
PRIOR FILING DATE: 2003-11-23
PRIOR PLING DATE: 2003-05-23
PRIOR APPLICATION NUMBER: PCT/US03/05346
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR PRILING DATE: 2003-02-20
PRIOR PILING DATE: 2003-02-20
PRIOR FILING DATE: 2003-02-30
PRIOR FILING DATE: 2003-
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NAME/KEY: misc_feature

LOCATION: (21)...(21)

OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-613
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OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match

4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 73.7%; Pred. No. 1e+02;
Matches 14; Conservative 5; Mismatches 0; Indels
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OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     148 CCACCGTTCATTCTAGAGC 166
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ORGANISM: Artificial Sequence
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NAME/KEY: misc feature
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LOCATION: (1)...(1)
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McWadgen, James
APPLICANT: McWadgen, James
APPLICANT: McWadgen, James
APPLICANT: McWadgen, James
TITLE OF INVENTION: RAY Interference Mediated Inhibition of Telomerase Gene
FILE OF INVENTION: Expression Using Short Interfering RNA (sinA)
FILE OF INVENTION: Expression Using Short Interfering RNA (sinA)
FILE OF INVENTION WUMBER: US/10/203,330
CURRENT PELLING DATE: 2004-08-20
CURRENT PELLING DATE: 2004-08-20
FRIOR PELLOR DATE: 2005-74
FRIOR PELLOR DATE: 2005-74
FRIOR PELLOR DATE: 2005-74
FRIOR PELLOR DATE: 2006-01-6
FRIOR PELLOR DATE: 2006-01-6
FRIOR FILING DATE: 2006-01-16
FRIOR FILING DATE: 2006-01-16
FRIOR FILING DATE: 2006-01-16
FRIOR FILING DATE: 2006-01-16
FRIOR FILING DATE: 2006-01-6
FRIOR FILING DATE: 2006-01-6
FRIOR FILING DATE: 2006-01-6
FRIOR FILING DATE: 2003-02-20
FRIOR FILING DATE: 2003-03-20
FRIOR FILING DATE: 2003-03-30
FRIOR FI
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PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR FILING DATE: 2003-02-20
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 768
SOFTWARE: Patentin version 3.3
SEQ ID NO 614
LENGTH: 21
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OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; LOCATION: (21) ... (21)
; OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-614
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4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 73.7%; Pred. No. 1e+02;
Matches 14; Conservative 5; Mismatches 0; Indels
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FEATURE:

LOCATION: (20) (20)

OTHER INFORMATION: n stands for thymidine BEATURE:

LOCATION: (21) (20)

OTHER INFORMATION: n stands for thymidine BEATURE:

LOCATION: (21) (21) (21)
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US-10-923-330-615
Sequence 615, Application US/10923330
Publication No. US20050153916A1
GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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US-10-923-330-618
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FILE REFERENCE: 400/209 (MBRBR: US/10/923,330

CURRENT APPLICATION NUMBER: US/10/923,330

CURRENT PELING DATE: 2004-08-24

PRIOR APPLICATION NUMBER: US 60/396,600

PRIOR FILING DATE: 2004-05-24

PRIOR APPLICATION NUMBER: US 10/826,966

PRIOR FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-01-14

PRIOR PELING DATE: 2004-01-14

PRIOR PELING DATE: 2004-01-14

PRIOR PELING DATE: 2003-01-14

PRIOR PELING DATE: 2003-01-14

PRIOR PELING DATE: 2003-01-14

PRIOR PELING DATE: 2003-01-23

PRIOR PELING DATE: 2003-01-23

PRIOR PELING DATE: 2003-05-23

PRIOR PELING DATE: 2003-05-23

PRIOR APPLICATION NUMBER: US 10/44,853

PRIOR APPLICATION NUMBER: US 10/44,853

PRIOR APPLICATION NUMBER: US 10/303/0508

PRIOR PELING DATE: 2003-02-20

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                                                                         OTHER INFORMATION: Description of Artificial Sequence: sinA sense region
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                                                                                                    FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(1)
OTHER:
NAME/KEY: misc_feature
LOCATION: (20)...(20)
OTHER:
NAME/KEY: misc_feature
LOCATION: (20)...(20)
OTHER:
NEORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: misc feature
LOCATION: (21) ... (21)
OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LOCATION: (1). (1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 19; DB 1; Length 21;
Pred. No. 1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Sequence 616, Application US/10923330; Publication No. US20050153916A1; GENERAL INPORMATION: APPLICANT: Sirna Therapeutics, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        150 ACCGTTCATTCTAGAGCAA 168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ACCGUCCAUCCUAGAGCAA 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 73.7%;
Matches 14; Conservative
ORGANISM: Artificial Sequence
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LOCATION: (1)..(1)
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT:
MCSWIGGEN, James
APPLICANT:
APPLICANT:
Beigelman, Leoning
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA INTERF. 2004-08-20
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT FILING DATE: 2004-08-20
FRIOR PILING DATE: 2004-06-24
FRIOR PILING DATE: 2004-05-24
FRIOR PELING DATE: 2004-06-24
FRIOR PELING DATE: 2004-01-17
FRIOR PILING DATE: 2004-01-14
FRIOR PILING DATE: 2004-01-14
FRIOR PILING DATE: 2004-01-14
FRIOR PILING DATE: 2003-01-24
FRIOR PILING DATE: 2003-01-24
FRIOR PILING DATE: 2003-01-24
FRIOR PILING DATE: 2003-01-34
FRIOR PILING DATE: 2003-02-20
FRIOR APPLICATION NUMBER: US 10/693,059
FRIOR PILING DATE: 2003-02-20
FRIOR APPLICATION NUMBER: US 10/503/05346
FRIOR APPLICATION NUMBER: US 10/503/05028
FRIOR APPLICATION NUMBER: US 2003-02-20
FRIOR PILING DATE: 2003-03-2-3
FRIOR FILING DATE: 2003-03-3-3
FRIOR FILING DATE: 2003-03-3-3
FRIOR FILING PAPEICATION NUMBER: PCT/US03/05028
FRIOR PLING PAPEICATION HORDER: PCT/US03/0
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OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
NAME/KEY: misc_feature

CCATION: (20)...(20)

COTHER INFORMATION: n stands for thymidine

FEATURE:

NAME/KEY: misc_feature

LOCATION: (21)...(21)

COTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-616
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                                                                                                                                                                                                                                                                                                                                                            Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                            Query Match
4.2%; Score 19; DB 1;
Best Local Similarity 73.7%; Pred. No. 1e+02;
Matches 14; Conservative 5; Mismatches
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OTHER INFORMATION: n stands for thymidine
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Publication No. US20050153916A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              300 GAAGAGTTGGGCTCTGTCA 318
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SEQ ID NO 618
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LOCATION: (21)..(21)
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APPLICANT: Beigelman, James
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RAD Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
CURRENT APPLICATION NUMBER: DC7/US03/0408
PRIOR PILING DATE: 2004-05-24
PRIOR PELING DATE: 2004-05-24
PRIOR PELING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-04-16
PRIOR FILING DATE: 2004-04-16
PRIOR PLING DATE: 2004-01-14
PRIOR PLING DATE: 2003-11-24
PRIOR PLING DATE: 2003-11-23
PRIOR APPLICATION NUMBER: US 10/693,059
PRIOR PLING DATE: 2003-02-20
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SOFTWARE: Patentin version 3.3
LENGTH: 21
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NAME/KEY: misc_feature

LOCATION: (21) - (21)

OTHER INFORMATION: n stands for thymidine
US-10-923-330-619
                                                                                                                                                                                                       Sequence 619, Application US/10923330
Publication No. US20050153916A1
GENERAL INFORMATION:
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US-10-923-330-620/c
Sequence 620, Application US/10923330
Publication No. US20050153916A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
145 CTTCCACCGTTCATTCTAG 163
                                     19 CTTCCACCGTTCATTCTAG 1
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ORGANISM: Artificial Sequence
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LOCATION: (20)..(20)
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ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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PRIOR FILING DATE: 2004-05-24
PRIOR PLICATION NUMBER: US 10/826,966
PRIOR FILING DATE: 2004-01-16
PRIOR FILING DATE: 2004-01-14
PRIOR PLICATION NUMBER: US 10/757,803
PRIOR PLICATION NUMBER: US 10/720,448
PRIOR PLICATION NUMBER: US 10/720,448
PRIOR FILING DATE: 2003-11-23
PRIOR FILING DATE: 2003-11-23
PRIOR FILING DATE: 2003-11-23
PRIOR PILING DATE: 2003-05-23
PRIOR PILING DATE: 2003-05-20
PRIOR PLICATION NUMBER: PCT/US03/05346
PRIOR FILING DATE: 2003-02-20
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COATION: (20)...(20)

COTER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
FRATURE:
NAME/KEY: misc_feature
LOCATION: (21)...(21)

COTHER INFORMATION: 010...(21)

US-10-923-330-621
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100.0%; Pred. No. 1e+02;
tive 0; Mismatches 0; Indels
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ORGANISM: Artificial Sequence
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Matches 19; Conservative
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US-10-923-330-622/c
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FEATURE: OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
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APPLICANT: Sirma Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leone
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
CURRENT APPLICATION NUMBER: US/10/923,330
PRIOR APPLICATION NUMBER: PCT/USO3/04088
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PRIOR PELLACATION NUMBER: PC1/0503/04088
PRIOR PELLOR DATE: 2004-05-24
PRIOR APPLICATION NUMBER: US 60/396,600
PRIOR PILING DATE: 2004-05-24
PRIOR PILING DATE: 2004-05-24
PRIOR PILING DATE: 2004-04-14
PRIOR PELLOATION NUMBER: US 10/75/803
PRIOR APPLICATION NUMBER: US 10/75/48
PRIOR APPLICATION NUMBER: US 10/720,448
PRIOR PILING DATE: 2003-11-23
PRIOR PILING DATE: 2003-11-23
PRIOR PELING DATE: 2003-11-23
PRIOR PELING DATE: 2003-12-20
PRIOR PELING DATE: 2003-02-20
PRIOR PELING DATE: 2003-02-20
PRIOR PILING DATE: 2003-02-20
PRIOR PELING DATE: 2003-02-20
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PRIOR FILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR FILING DATE: 2003-02-20
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ. ID NOS: 768
SOFTWARE: PatentIn version 3.3
SQI ID NO 622
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LOCATION: (20) ... (20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
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1e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
NAME/KEY: misc_feature
LOCATION: (21) - (21)
UCHER INFORMATION: n stands for thymidine
US-10-923-330-622
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Conservative
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SEQUENCE 624, Application US/10923330

Publication No. US20050153916A1

GENERAL INFORMATION:
Publication No. US20050153916A1

GENERAL INFORMATION:
Publication No. US20050153916A1

APPLICANT: Beigelman, Leonid

TITLE OF INVENTION: RAPESSON USING SINCE Interfering RNA (siNA)

TITLE OF INVENTION: RAPESSON USING SINCE Interfering RNA (siNA)

TITLE OF INVENTION NUMBER: PCT/US03/04088

PRIOR APPLICATION NUMBER: US 60/396,600

PRIOR PELING DATE: 2004-05-24

PRIOR PELING DATE: 2002-07-17

PRIOR PELING DATE: 2002-07-17

PRIOR PELING DATE: 2002-07-17

PRIOR PELING DATE: 2004-04-16

PRIOR PELING DATE: 2004-04-16

PRIOR APPLICATION NUMBER: US 10/757,803

PRIOR APPLICATION NUMBER: US 10/757,803

PRIOR PELING DATE: 2003-11-23

PRIOR PELING DATE: 2003-12-20

PRIOR PELING DATE: 2003-11-23

PRIOR PELING DATE: 2003-12-20

PRIOR PELING DATE: 2003-11-33

PRIOR PELING DATE: 2003-02-20

PRIOR PELING DAT
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LOCATION: (21)..(21)

// OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage

US-10-923-330-623
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LOCATION: (20)_.(20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
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hes 0; Indels
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                                                                                                                                                                                                                                                                                                                                             Query Match 4.2%; Score 19; DB Best Local Similarity 100.0%; Pred. No. le-Matches 19; Conservative 0; Mismatches
FEATURE:
NAME/KEY: misc_feature
LOCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
FEATURE:
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; OTHER INFORMATION: n stands for thymidine
US-10-923-330-624
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Best Local Similarity
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Sequence 626, Application US/10923330
| Sequence 626, Application US/10923330
| Publication No. US20050153916A1
| GENERAL INFORMATION:
| APPLICANT: Sitna Therapeutics, Inc.
| APPLICANT: MoSWAGEN, James
| APPLICANT: MOSWAGEN, James
| APPLICANT: MOSWAGEN, James
| APPLICANT: MOSWAGEN, James
| APPLICANT: Beigelman, Leonid
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
| FILE REFERENCE: 400/209 (MRH02-708-C)
| CURRENT PAPLICATION NUMBER: US/10/923,330
| CURRENT PELLOR DATE: 2004-06-20
| PRIOR PILING DATE: 2004-06-20
| PRIOR PELLOR DATE: 2004-06-20
| PRIOR PELLOR DATE: 2004-06-20
| PRIOR PELLOR DATE: 2004-06-10
| PRIOR PELLOR DATE: 2004-06-10
| PRIOR PELLOR DATE: 2004-06-10
| PRIOR PELLOR DATE: 2004-01-11-24
| PRIOR PELLOR DATE: 2004-01-12
| PRIOR PELLOR DATE: 2003-11-23
| PRIOR PELLOR DATE: 2003-05-30
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        Gaps
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0
        Indels
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0; Mismatches
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NAME/KEY: misc_feature
LOCATION: (1)._(2)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: misc_feature
LOCATION: (7)._(7)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LOCATION: (11)...(11)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                          19 GAAGAGITGGGCTCTGTCA 1
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LOCATION: (3)...(6)
OTHER INFORMATION: 2'-O-methyl
FEATURE:
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LOCATION: (8)..(10)
OTHER INFORMATION: 2'-O-methyl
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LOCATION: (12)...(13)
OTHER INFORMATION: 2'-O-methyl
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ORGANISM: Artificial Sequence
FEATURE:
                                                                 300 GAAGAGTTGGGCTCT
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US-10-923-330-628/c
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                                                                                                                                                                                       Gaps
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                                                                                                                  Score 19; DB 1; Length 21;
Pred. No. 1e+02;
0; Mismatches 0; Indels
                                                                                                                                                                 Query Match 4.2%; Score 19; DB Best Local Similarity 100.0%; Pred. No. 1e+Matches 19; Conservative 0; Mismatches
                                                                 FEATURE:
NAME/KEY: misc feature
LOCATION: (20) ... (20)
OTHER INFORMATION: n stands for thymidine
NAME/KEY: misc_feature
LOCATION: (14) ... (14)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
NAME/KEX: misc_feature
LOCATION: (1). (3)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                          145 CTTCCACCGTTCATTCTAG 163
                                                                                                                                                                                                                           19 CTTCCACCGTTCATTCTAG 1
                          FEATURE:
NAME/KEY: misc_feature
LOCATION: (15)..(19)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                    US-10-923-330-627/c
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US-10-924-310-628(Application US/10923330)

| US-10-924-310-628(Application US/10923330)
| Publication No. US20050153916A1 |
| GENERAL INFORMATION:
| APPLICANT: Sirna Therapoutics, Inc. |
| APPLICANT: Beigelman, James |
| APPLICANT: Beigelman, Leonid |
| TITLE OF INVENTION: Expression Using Short Interfering RNA (BiNA) |
| TITLE OF INVENTION: Expression Using Short Interfering RNA (BiNA) |
| TITLE OF INVENTION: Expression Using Short Interfering RNA (BiNA) |
| TITLE OF INVENTION: Expression Using Short Interfering RNA (BiNA) |
| TITLE OF INVENTION: Expression Using Short Interfering RNA (BiNA) |
| TITLE OF INVENTION: Expression Using Short Interfering RNA (BiNA) |
| FRIOR FILING DATE: 2004-08-20 |
| PRIOR FILING DATE: 2004-05-24 |
| PRIOR FILING DATE: 2004-05-24 |
| PRIOR PELICATION NUMBER: US 10/826,966 |
| PRIOR APPLICATION NUMBER: US 10/757,803 |
| PRIOR FILING DATE: 2004-04-16 |
| PRIOR FILING DATE: 2004-04-16 |
| PRIOR FILING DATE: 2004-01-14 |
| PRIOR FILING DATE: 2003-11-24 |
| PRIOR FILING DATE: 2004-01-14 |
| PRIOR FILING DATE: 2003-11-24 |
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OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                   NAME/KEY: misc feature
LOCATION: (12) ... (12)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                   LOCATION: (8). (8)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LOCATION: (15)..(15)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (4).7(7)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LOCATION: (16)..(19)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                         NAME/KEY: misc_feature
LOCATION: (9)._(11)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LOCATION: (13)...(14)
OTHER INFORMATION: 2'-0-methyl
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Best Local Similarity 100.0
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: misc_feature
                                                                                   NAME/KEY: misc_feature
LOCATION: (8)..(8)
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FEATURE:
NAME/KEY: misc_feature
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PRIOR FILING DATE: 2003-11-23
PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR FILING DATE: 2003-05-23
PRIOR FILING DATE: 2003-05-20
PRIOR FILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR PILING DATE: 2003-02-20
NUMBER OF SEQ ID NOS: 768
SOFTWARE: Patentin version 3.3
SEQ ID NO 628
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; LOCATION: (21)_..(21)
; CTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-628
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4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
NAME/KEY: misc_feature
LOCATION: (5)...(8)
OTHER INFORMATION: 2'-0-methyl
FEATURE:
NAME/KEY: misc_feature
LOCATION: (9)...(9)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: misc feature
LOCATION: (20) ... (20)
OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                                                                    LOCATION: (1)..(4)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LOCATION: (13)...(13)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: misc_feature
LOCATION: (16)...(16)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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US-10-923-330-629/c
Sequence 629, Application US/10923330
Publication No. US20050153916A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            147 TCCACGTTCATTCTAGAG 165
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LOCATION: (10)...(12)
OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: misc_feature
LOCATION: (14)...(15)
OTHER INFORMATION: 2'-O-methyl
                                                                                                                                                                                                                                                                         TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LOCATION: (17)...(19)
OTHER INFORMATION: 2'-O-methyl
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LOCATION: (1)..(4)
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LOCATION: (13)..(13)
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LOCATION: (17)..(19)
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APPLICANT: Resergeen, James
TITLE OF INVESTIONS PRACTICES AND THE SERVICES AND THE SERVICES
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US-10-292-330-630.

US-10-292-330-630.

US-10-292-330-630.

Publication No. US20050153916A1

GENERAL INFORMATION

MEDICANT: MCSAVIGGEN, James

APPLICANT: MCSAVIGGEN, James

TITLE OF INVENTION: REPTERENCE: 400/209 (MBHE02-708-C)

CURRENT PLING DATE: 2004-05-24

PRIOR PAPLICATION NUMBER: DCT/US03/0408

PRIOR PLING DATE: 2004-05-24

PRIOR APPLICATION NUMBER: US 60/396,600

PRIOR APPLICATION NUMBER: US 60/396,600

PRIOR PRIOR DATE: 2004-05-24

PRIOR APPLICATION NUMBER: US 10/720,966

PRIOR PLING DATE: 2004-04-16

PRIOR PLING DATE: 2004-04-16

PRIOR PLING DATE: 2004-01-14

PRIOR PLING DATE: 2003-01-17

PRIOR PLING DATE: 2003-01-12

PRIOR PLING DATE: 2003-01-12

PRIOR PLING DATE: 2003-01-24

PRIOR PLING DATE: 2003-01-34

PRIOR PLING
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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NAME/KEY: misc_feature
LOCATION: (20)
COCATION: (20)
POTHER INFORMATION: n stands for thymidine
NAME/KEY: misc_feature
LOCATION: (21)...(21)
COTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-629
                                                                                                                                                                                                                                                                                                                                                    Query Match

4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches 0; Indels
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LOCATION: (3)..(6)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
FEATURE:
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LOCATION: (1)._(1)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      148 CCACCGTTCATTCTAGAGC 166
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LOCATION: (2)...(2)
OTHER INFORMATION: 2'-0-methyl
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 211
US-10-923-330-630/c
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GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RAN Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RAN Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: RAN INTERS.
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REFERENCE: 400/209 (MBHBO2-708-C)
CURRENT APPLICATION NUMBER: PCT/US03/04088
PRIOR PILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-24
PRIOR APPLICATION NUMBER: US 10/826,966
PRIOR FILING DATE: 2004-04-16
PRIOR PRILING DATE: 2004-04-16
PRIOR PRILING DATE: 2004-01-14
PRIOR PRILING DATE: 2004-01-14
PRIOR PRILING DATE: 2004-01-14
PRIOR FILING DATE: 2004-01-14
PRIOR FILING DATE: 2003-01-124
PRIOR FILING DATE: 2003-11-24
PRIOR FILING DATE: 2003-11-24
PRIOR FILING DATE: 2003-11-23
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; OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-630
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 100.0%; Pred. No. 1e+02; Matches 19; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: misc feature
LOCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
                                                                                                     NAME/KEY: misc feature
LOCATION: (11) ... (11)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                               LOCATION: (15)...(15)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc feature
LOCATION: (18)...(18)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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; Sequence 631, Application US/10923330
; Publication No. US20050153916A1
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                                                                                                                                                                                                                           NAME/KEY: misc_feature
LOCATION: (12)...(14)
OTHER INFORMATION: 2'-O-methyl
                                                 OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: misc_feature
LOCATION: (16)..(17)
OTHER INFORMATION: 2'-O-methyl
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LOCATION: (19)..(19)
OTHER INFORMATION: 2'-O-methyl
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NAME/KEY: misc feature
NAME/KEY: misc_feature
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Publication No. US20050153916A1
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PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR FILING DATE: 2003-05-23
PRIOR PPLICATION NUMBER: PCT/US03/05346
PRIOR PLIING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: PCT/US03/05028
PRIOR FILING DATE: 2003-02-20
PRIOR FILING PAID: 2003-02-20
Remaining Prior Application data removed - See File Wrapper or PALM.
SOFTWARE: Patentin version 3.3
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NAME/KEY: misc_feature
NAME/KEY: (21)..(21)
OCHER_INFORMATION: 3'-3 attached terminal deoxyabasic moeity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 100.0%; Pred. No. 1e+02; Matches 19; Conservative 0; Mismatches 0; Indels
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NAME/KEY: misc_feature
LOCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: misc_feature
LOCATION: (1). (2)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: misc_feature
LOCATION: (4)._(7)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (12)...(12)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (16)...(16)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY:
MACATION: (19)...(19)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (3)...(3)
OTHER INFORMATION: 2'-O-methyl
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NAME/KRY: misc feature
LOCATION: (8). (1).

OTHER INFORMATION: 2'-0-methyl
                                                                                                                                                                                                                                                        TYPE: RNA
ORGANISM: Artificial Sequence
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OTHER INFORMATION: 2'-0-methyl
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OTHER INFORMATION: 2'-0-methyl
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LOCATION: (17)..(18)
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                                                                                                                                                                                                            SEQ ID NO 631
LENGTH: 21
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RESULT 213 US-10-923-330-632/c ; Sequence 632, Application US/10923330

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APPLICANT: Mos. 1920.0513316A1

TITLE OF INVENTION: RAN INTERference Modiated Inhibition of Polomerase Gene IIILE OF INVENTION: RAN INTERference Modiated Inhibition of Polomerase Gene IIILE OF INVENTION: RAN INTERference Modiated Inhibition of Polomerase Gene IIILE OF INVENTION: RAN INTERFERENCE ADDISONAL MORRES. PCT (1920.704-0)

PRIOR PELING DATE: 2004-00-17

PRIOR FILING DATE: 2004-00-17

PRIOR FILING DATE: 2004-00-17

PRIOR FILING DATE: 2004-00-17

PRIOR FILING DATE: 2004-00-17

PRIOR PELING DATE: 2004-00-14

PRIOR PERING DATE: 2004-00-14

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                                                                                                                                OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
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; LOCATION: (21)...(21); OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity US-10-923-330-632
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; OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-634
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                                          4.2%; Score 19; DB 1; Length 21;
100.0%; Pred. No. 1e+02;
tive 0; Mismatches 0; Indels
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                                                                                   300 GAAGAGTTGGGCTCTGTCA 318
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                                                                                               19 GAAGAGTTGGGCTCTGTCA 1
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ORGANISM: Artificial Sequence
                                       Query Match 4.2%;
Best Local Similarity 100.0%
Matches 19; Conservative
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Sequence 638, Application US/10923330

Sequence 638, Application US/10923330

Sequence 638, Application US/10923330

Sequence 638, Application Worksygon, James

APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Beigelman, Leonid

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)

CURRENT APPLICATION NUMBER: US/10/923,330

CURRENT APPLICATION NUMBER: PCT/US03/04088

PRIOR APPLICATION NUMBER: D04-00-20

PRIOR APPLICATION NUMBER: D10/04088

PRIOR APPLICATION NUMBER: US 60/396,600

PRIOR APPLICATION NUMBER: US 10/926,400

PRIOR FILING DATE: 2002-07-17

PRIOR PLING DATE: 2003-05-24

PRIOR PLING DATE: 2003-01-14

PRIOR APPLICATION NUMBER: US 10/757,803

PRIOR PLING DATE: 2003-11-23

PRIOR APPLICATION NUMBER: US 10/757,803

PRIOR APPLICATION NUMBER: US 10/693,059

PRIOR APPLICATION NUMBER: US 10/693,059

PRIOR APPLICATION NUMBER: US 10/644,853

PRIOR APPLICATION NUMBER: PCT/US03/05346

PRIOR PLING DATE: 2003-02-23

PRIOR APPLICATION NUMBER: PCT/US03/05346

PRIOR PLING DATE: 2003-02-23

PRIOR APPLICATION NUMBER: PCT/US03/05346

PRIOR APPLICATION NUMBER: PCT/US03/05346

PRIOR APPLICATION NUMBER: PCT/US03/05346

PRIOR PLING DATE: 2003-02-23

PRIOR PLING DATE: 2003-02-23

PRIOR APPLICATION NUMBER: PCT/US03/05346

PRIOR APPLICATION NUMBER: PCT/US03/05346

PRIOR PLING DATE: 2003-02-23

PRIOR PLING DATE: 2003-02-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
                                                              PRIOR FILING DATE: 2003-11-24

PRIOR PLING DATE: 2003-11-23

PRIOR APPLICATION NUMBER: US 10/693,059

PRIOR APPLICATION NUMBER: US 10/444,853

PRIOR PILING DATE: 2003-05-20

PRIOR FILING DATE: 2003-02-20

PRIOR PILING DATE: 2003-02-20

PRIOR PILING DATE: 2003-02-20

PRIOR PILING DATE: 2003-02-20

Remaining Prior Application data removed - See File Wrapper or PALM.

NUMBER OF SEQ ID NOS: 768

SOFTWARE: Patentin version 3.3

LENGTH: 21
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NUMBER OF SEQ ID NOS: 768
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NAME/KEY: misc feature

LOCATION: (20)

CTHER INFORMATION: n stands for thymidine

FEATURE:

NAME/KEY: misc feature

LOCATION: (21)

CTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity

US-10-923-330-637
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100.0%; Pred. No. ...
... 0; Mismatches
                                                 US 10/720,448
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Best Local Similarity 100.0
Matches 19; Conservative
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| Publication No. US20050153916A1
| GENERAL INFORMATION:
| APPLICANT: Strna Therapeutics, Inc.
| APPLICANT: Strna Therapeutics, Inc.
| APPLICANT: Beigaham, Leonid
| TITLE OF INVENTION: Expression Using Short Interfering RNA (sinA)
| TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
| TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
| TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase
| TITLE OF INVENTION: RNA INTERFERENCE: 400/209 (MEHBO2-708-C)
| FILE REFERENCE: 400/209 (MEHBO2-708-C)
| FRICH REPLICATION NUMBER: PCT/US03/04088
| PRIOR FILING DATE: 2004-06-24
| PRIOR PLICATION NUMBER: US 60/396,600
| PRIOR PLICATION NUMBER: US 10/426,566
| PRIOR FILING DATE: 2004-04-16-24
| PRIOR FILING DATE: 2004-04-16-14
| PRIOR FILING DATE: 2004-04-16
| PRIOR FILING DATE: 2004-04-16
| PRIOR FILING DATE: 2004-04-16
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                                 CURRENT FILING DATE: 2004-08-20

PRIOR PELING DATE: 2004-08-20

PRIOR PELING DATE: 2004-06-24

PRIOR FILING DATE: 2004-06-24

PRIOR PELING DATE: 2004-06-136

PRIOR PELING DATE: 2002-07-17

PRIOR APPLICATION NUMBER: US 10/826,966

PRIOR PILING DATE: 2004-04-16

PRIOR PILING DATE: 2004-04-14

PRIOR PILING DATE: 2004-01-14

PRIOR PILING DATE: 2003-11-23

PRIOR APPLICATION NUMBER: US 10/720,448

PRIOR APPLICATION NUMBER: US 10/693,059

PRIOR PILING DATE: 2003-11-23

PRIOR PILING DATE: 2003-11-23

PRIOR PILING DATE: 2003-12-23

PRIOR PILING DATE: 2003-02-20

PRIOR APPLICATION NUMBER: PCT/US03/05028

PRIOR PILING DATE: 2003-02-20

PRIOR PILING DATE: 2003-02-20
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; OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity US-10-923-330-636
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4.2%; Score 19; DB 1; Length 21;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches 0; Indels
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LOCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
CURRENT APPLICATION NUMBER: US/10/923,330
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ORGANISM: Artificial Sequence
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LENGTH: 21
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RESULT 220
US-10-923-330-640/c
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Sequence 639, Application Us/10923330

Publication No. US20050153916A1

APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: Beigelman, Leonid

TITLE OF INVENTION: RAN Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RAN Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: RAN Interference Mediated Inhibition of Telomerase Gene

TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)

FILE REFERENCE: 400/209 (MBHB02-708-C)

CURRENT FILING DATE: 2004-05-24

PRIOR FILING DATE: 2004-05-24

PRIOR PELING DATE: 2004-05-24

PRIOR PELING DATE: 2004-01-14

PRIOR APPLICATION NUMBER: US 10/856,966

PRIOR APPLICATION NUMBER: US 10/757,803

PRIOR PILING DATE: 2003-11-24

PRIOR PILING DATE: 2003-10-24

PRIOR PILING DATE: 2003-10-3

PRIOR PILING DATE: 2003-10-3

PRIOR PILING DATE: 2003-02-20

PRIOR PILING DATE
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; OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-638
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 4.2%; Score 19; DB 1; Length 21; Best Local Similarity 100.0%; Pred. No. 1e+02; Matches 19; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                       FEATURE:
NAME/KEY: misc_feature
LOCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
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LOCATION: (20)..(20)
OTHER INFORMATION: n stands for thymidine
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         149 CACCGTTCATTCTAGAGCA 167
SOFTWARE: PatentIn version 3.3
SEQ ID NO 638
LENGTH: 21
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ORGANISM: Artificial Sequence
                                                                                                                TYPE: RNA
ORGANISM: Artificial Sequence
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US-10-923-330-639/c
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GENERAL INFORMATION:
JAPPLICANT: Sirna Therapeutice, Inc.
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Telomerase Gene
TITLE OF INVENTION: Expression Using Short Interfering RNA (siNA)
FILE REPRENCE: 400/209 (MBH03C-708-C)
CURRENT APPLICATION NUMBER: US/10/923,330
CURRENT APPLICATION NUMBER: PCT/US03/0408
PRIOR FILING DATE: 2004-06-20
PRIOR PILING DATE: 2004-06-24
PRIOR FILING DATE: 2004-06-24
PRIOR FILING DATE: 2004-07-17
PRIOR PELICATION NUMBER: US 10/757,803
PRIOR FILING DATE: 2004-01-14
PRIOR FILING DATE: 2004-01-14
PRIOR FILING DATE: 2004-01-14
PRIOR FILING DATE: 2004-01-14
PRIOR FILING DATE: 2004-01-23
PRIOR PELICATION NUMBER: US 10/693,059
PRIOR FILING DATE: 2003-01-24
PRIOR FILING DATE: 2003-01-24
PRIOR PELICATION NUMBER: US 10/444,853
PRIOR PELICATION NUMBER: PCT/US03/05028
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                                                                                                                                                                                                                                                                                   Gарв
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; NAME/KEY: misc_feature
; LOCATION: (21)...(21)
; OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-923-330-639
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ) NAME/KEY: misc_feature

; LOCATION: (21):.(21)

; OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity

US-10-923-330-640
                                                                                                                                                                                                    Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 21;
                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                    ch 4.2%; Score 19; DB 1;
1 Similarity 100.0%; Pred. No. 1e+02;
19; Conservative 0; Mismatches (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
4.2%; Score 19; DB 1;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 19; Conservative 0; Mismatches C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: misc_feature
LCCATION: (20)...(20)
OTHER INFORMATION: n stands for thymidine
PEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 640, Application US/10923330 Publication No. US20050153916A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                   150 ACCGTTCATTCTAGAGCAA 168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         300 GAAGAGTTGGGCTCTGTCA 318
                                                                                                                                                                                                                                                                                                                                                                                                                 19 ACCGTTCATTCTAGAGCAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19 GAAGAGTIGGGCTCTGTCA 1
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                                                                                                                                                                                                                                         Best Local Similarity
Matches 19; Conserv
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FEATURE
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                                                                                  APPLICANT: Kilian, Andrzej
APPLICANT: Kilian, Andrzej
APPLICANT: Bowtell, David
TITLE OF INVENTION: VERTEBRATE TELOMERASE GENES AND PROTEINS AND USES
TITLE OF INVENTION: THEREOF
FILE REFERENCE: TWGL 40723
CURRENT APPLICATION NUMBER: US/10/900,231
CURRENT FILING DATE: 2004-07-27
NUMBER OF SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 126
LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: Description of Artificial Sequence: Synthesized OTHER INFORMATION: Amplification Primer Design based on EST Sequence; OTHER INFORMATION: GenBank Accession Number AA281296 US-10-900-231-126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .
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| Publication No. US20030204069A1
| GENERAL INFORMATION:
| Lingner, Joachim | Nakamura, Toru | Chapman, Karen B. | Morin, Gregg B. | Harley, Calvin B. | Andrews, William H. | TITLE OF INVENTION: William H. | ITTLE OF INVENTION: Human Telomerase Catalytic Subunit | NUMBER OF SEQUENCES: 633
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4.0%; Score 18; DB 1; Length 18;
100.0%; Pred. No. 93;
cive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/325,810
FILING DATE: 20-06-2002
CLASSIFICATION: «Unknown»
PRIOR APPLICATION NUMBER: US/09/402,181
FILING DATE: 29-56p-1997
APPLICATION NUMBER: US/09/402,181
APPLICATION NUMBER: US/09/402,481
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 01-CCT-1996
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/846,017
                      Sequence 126, Application US/10900231
Publication No. US20050176022A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 GGGTTGCGGAGGGTGGGC 18
                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 GGGTTGCGGAGGGTGGGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 100.
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 222
US-10-325-810-543/c
US-10-900-231-126
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Sequence 543, Application US/10877124

Publication No. US20040242529A1

GENERAL INPORMATION:

Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
Morin, Greeg B.

Morin, Greeg B.

Harley, Calvin B.

Andrews, William H.

TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 727

CORRESPONDENCE ADDRESS:

ADDRESSE: Townsend and Townsend and Crew Lip
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California

COUNTRY: USA

IN PAINI-1834

COMPUTER READBALE FORM:

MEDIUM TYPE: Floppy disk.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; LOCATION: 1..18
OTHER INFORMATION: /note= "antisense hTRT molecule"; SEQUENCE DESCRIPTION: SEQ ID NO: 543:
US-10-325-810-543
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3.6%; Score 16.4; DB 1; Length 1 94.4%; Pred. No. 1.48+02; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/877,124
FILING DATE: 24-Jun-2004
CLASSIFICATION: <UNKNOWN>
           APPLICATION NUMBER: US 08/851,843
                                                                                                                                                                                                                                                                                                               (415) 576-0200
                                                                                                                                                                                                                                                                                                                             TELEFAX: (415) 576-0300 INFORMATION FOR SEQ ID NO: 543: SEQUENCE CHARACTERISTICS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              149 CACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 18 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18 caccerrcarreragase 1
                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 3.6
Best Local Similarity 94.4
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                             TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY:
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US-10-877-124-543/c
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 3.6%; Score 16.4; DB 1; Length 18; Best Local Similarity 94.4%; Pred. No. 1.4e+02; Matches 17; Conservative 0; Mismatches 1; Indels
                                                                                                                         SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/877,022
FILING DATE: 24-Jun-2004
CLASSIFICATION: <UNKNOWN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: WO PCT/US97/17618
FILING DATE: 01-OCT-1997
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-OCT-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002610US
                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/09/432,503
FILING DATE: 02-Nov-1999
APPLICATION NUMBER: 08/974,549
FILING DATE: 4UNCOMN
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 09-MAY-1997
APPLICATION NUMBER: US 08/851,312
APPLICATION NUMBER: US 08/851,312
APPLICATION NUMBER: US 08/854,050
FILING DATE: 09-MAY-1997
APPLICATION NUMBER: US 08/854,050
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-877-146-543/c; Sequence 543, Application US/10877146; Publication No. US20050013825A1; GENERAL INPORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Gech, Thomas R.
Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        149 CACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 18 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (415) 576-03
INFORMATION FOR SEQ ID NO: 543:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18 caccerrcarreragase
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OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-10-877-022-543
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Morin, Garegi B.
Harley, Calvin B.
Harley, Calvin B.
Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 727
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LOCATION: 1..18
OTHER INFORMATION: /note= "antisense hTRT molecule"
SEQUENCE DESCRIPTION: SEQ ID NO: 543:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 3.6%; Score 16.4; DB 1; Length 18; Best Local Similarity 94.4%; Pred. No. 1.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002610US
TELECOMMUNICATION INFORMATION:
                APPLICATION NUMBER: US/09/432,503
FILING DATE: 02-Nov-1999
FILING DATE: 02-Nov-1999
FILING DATE: 02-Nov-1999
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/851,050
FILING DATE: 09-MAY-1997
APPLICATION NUMBER: US 08/911,312
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: WO PCT/US97/17618
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: WO PCT/US97/17618
FILING DATE: 01-OCT-1997
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-OCT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 224
US-10-877-022-543/c
iSequence 543, Application US/10877022
i Publication No. US20040247613A1
i GENERAL INFORMATION:
i APPLICANT: Cech, Thomas R.
i Lingner, Joachim
i Nakamura, Toru
i Chapman, Karen B.
Chapman, Karen B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 543:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         149 CACCGTTCATTCTAGAGC 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 18 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18 CACCCTTCATTCTAGAGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17; Conservative
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RESULT 226
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US-10-877-146-543
                                                                                                                                                                                                                                                                                  COMPUTER: USAA

ZIP: 94111-3834

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBW PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/877,146

CLASSIFICATION NUMBER: US/99/432,503

FILING DATE: CA-Nov-1999
APPLICATION NUMBER: US/99/42,419

FILING DATE: CHENOWN
APPLICATION NUMBER: US 08/844,419

FILING DATE: US-RAY-1997
APPLICATION NUMBER: US 08/846,017

FILING DATE: US-APR-1997
APPLICATION NUMBER: US 08/854,050

FILING DATE: US-APR-1997
APPLICATION NUMBER: US 08/911,312

FILING DATE: US-MAY-1997
APPLICATION NUMBER: US 08/911,312

FILING DATE: US-MAY-1997
APPLICATION NUMBER: US 08/911,312

FILING DATE: US-MAY-1997
APPLICATION NUMBER: US 08/915,503

FILING DATE: US-MC-1997
APPLICATION NUMBER: WO PCT/US97/17618

FILING DATE: US-MAY-1997
APPLICATION NUMBER: WO PCT/US97/17618

FILING DATE: US-MAY-1997
APPLICATION NUMBER: US 08/915,503

FILING DATE: US-MAY-1997
APPLICATION NUMBER: US 08/915,503

FILING DATE: US-MAY-1997
APPLICATION NUMBER: US 08/915,503

FILING DATE: US-MAY-1997
APPLICATION NUMBER: WO PCT/US97/17618

FILING DATE: US-MAY-1997
APPLICATION NUMBER: US 08/915,503
                            Andrews, William H.
DF INVENTION: Human Telomerase Catalytic Subunit
OF SEQUENCES: 727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 16.4; DB 1; Length 18;
Pred. No. 1.4e+02;
                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002610US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
Harley, Calvin B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: 543:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS
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OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear
MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3.69
Best Local Similarity 94.49
Matches 17, Conservative
                                                      TITLE OF NUMBER OF
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                                                                                            APPLICANT: Gorman, Cornelia M.,
Groskreutz, Debyra J.
TITLE OF INVENTION: Prohormone Convertase Transformed Cells and
Polypeptide Synthesis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-10-900-231-127/c
US-10-900-231-127/c
Fequence 127, Application US/10900231
Fequence 127, Application No. US20050176022A1
Fublication No. US2005017602A1
GENERAL INFORMATION:
APPLICANT: Mountain, Andrzej
APPLICANT: Moutell, David
TITLE OF INVENTION: THEREOF
TITLE OF INVENTION: THEREOF
TITLE OF INVENTION: THEREOF
CURRENT APPLICATION NUMBER: US/10/900,231
CURRENT FILING DATE: 2004-07-27
NUMBER OF SEQ ID NOS: 155
SOFTWARE: Patentin Ver. 2.0
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85.7%; Pred. No. 2e+02;
Live 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk COMPUTER: 11M PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/997,868
FILING DATE: 12-Mar-2002
CLASSIFICATION NUMBER: US/09/997,868
FILING DATE: 12-Mar-2002
CLASSIFICATION NUMBER: 07/887265
FILING DATE: 22-May-1992
APPLICATION NUMBER: 07/803631
FILING DATE: 06-DEC-1992
APPLICATION NUMBER: PCT/US92/10621
FILING DATE: 04-DEC-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Love, Richard B.
REGISTRATION NUMBER: 34,659
REFERENCE/DOCKET NUMBER: P0748P3
TELECOMPUNICATION INFORMATION:
TELEPHONE: 650/225-530
TELEFRAX: 650/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE DESCRIPTION: SEQ ID NO: 13:
                                                                                                                                                                                         NUMBER OF SEQUENCES: 57
CORRESPONDENCE ADDRESS:
ADDRESSE:
STREET: 1 DNA WAY
CITY: South San Francisco
STATE: California
COUNTRY: USA
                         Sequence 13, Application US/09997868; Publication No. US20030031654A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   156 CATTCTAGAGCAAAAAAA 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CATTCTAGAGCAAAGAGACAA 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                           ZIP: 94080
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18; Conservative
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US-09-997-868-13
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Matches
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Gaps

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Indels

0; Mismatches

149 CACCGTTCATTCTAGAGC 166

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18 CACCCTTCATTCTAGAGC 1

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Sequence 76, Application US/10071179

Publication No. US20030108862A1

GENERAL INFORMATION:
ITLE OF INVENTION: A NUCLEIC ACID ENCODING A RETINOBLASTOMA BINDING PROTEIN (RBP-7)

TITLE OF INVENTION: AND POLYMORPHIC MARKERS ASSOCIATED WITH SAID NUCLEIC ACID.

TITLE OF INVENTION: AND POLYMORPHIC MARKERS ASSOCIATED WITH SAID NUCLEIC ACID.

FILE REFERENCE: GENSET. 031A

CURRENT APPLICATION NUMBER: US/10/071,179

CURRENT APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/345,882

PRIOR PILING DATE: EARLIER PILING DATE: 1999-06-30

PRIOR FILING DATE: EARLIER PILING DATE: 1998-06-30

PRIOR FILING DATE: EARLIER PILING DATE: 1998-06-10

PRIOR FILING DATE: EARLIER FILING DATE: 1998-12-10

NUMBER OF SEQ ID NOS: 140

SOFTWARE: PATENT.

SEQ ID NOS: 140

SEC ID NOS: 140
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LOCATION: 1..20
COTHER INFORMATION: upstream amplification primer for SEQ 34, SEQ 55, SEQ 35, SEQ 56
US-10-071-179-76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
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                                                                             Sequence 84, Application US/10714195
Sequence 84, Application US/10714195
Sequence 84, Application US/001978541
SERBEAL INFORMATION:
APPLICANT: BARE, Joffre
APPLICANT: Cronin, Maureen
APPLICANT: BAR, Steve
APPLICANT: BAR, Steve
TITLE OF INVENTION: GENE EXPRESSION PROFILING OF EGFR
TITLE OF INVENTION: POSITIVE CANCER
FILE REFERENCE: 39740-0005
CURRENT FILING DATE: 2003-11-14
PRIOR APPLICATION NUMBER: 60/427090
PRIOR APPLICATION NUMBER: 60/427090
PRIOR APPLICATION NUMBER: 60/427090
SRIOR FILING DATE: 2003-11-15
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 15.4; DB 1;
Pred. No. 2.1e+02;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           166 CAAACAAAAATGTCAG 182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 CAAACAATAAATGTCAG 17
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 3.4%;
Best Local Similarity 94.1%;
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 54.2%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 231
US-10-071-179-76
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 71, Application US/10174020
Publication No. US20030232770A1
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
TAPLICANT: Breat P. Monia
APPLICANT: Kenneth W. Dobie
TITLE OF INVENTION: ANTISENSE MODULATION OF HYPOTHETICAL TUMOR ENDOTHELIAL MARKER EXF
FILE REFERENCE: RTS-03869
CURRENT APPLICATION NUMBER: US/10/174,020
CURRENT FILING DATE: 2002-06-17
NUMBER OF SEQ ID NOS: 149
SEQ ID NO 71
LENGTH: 20
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Publication No. US20030232770A1
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Brett P. Monia
APPLICANT: Renneth W. Dobie
TITLE OF INVERTION: ANTISENSE MODULATION OF HYPOTHETICAL TUMOR ENDOTHELIAL MARKER EXF
FILE REFERENCE: RTS-0369
CURRENT APPLICATION NUMBER: US/10/174,020
CURRENT FILING DATE: 2002-06-17
NUMBER OF SEQ ID NOS: 149
SEQ ID NO 138
LENGTH: 20
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                                                                                                OTHER INFORMATION: Description of Artificial Sequence: Synthesized OTHER INFORMATION: Amplification Primer Design based on EST Sequence; OTHER INFORMATION: GenBank Accession Number AA281296
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
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                                                                                                                                                                                                                                                                          Query Match 3.6%; Score 16.2; DB 1; Length 21; Best Local Similarity 85.7%; Pred. No. 2e+02; Matches 18; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3.5%; Score 15.8; DB 1; Length 20; 89.5%; Pred. No. 1.9e+02; Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 15.8; DB 1; Length 20; Pred. No. 1.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: Antisense Oligonucleotide
US-10-174-020-71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                          431 CAGGACTCGGCTCACACATGC 451
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21 CAGGACTCGGCTCACCACTGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    362 GGCCGCAGGAAGAGAACG 380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 362 GGCCGCAGGAAGGAACG 380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 GGCCACAGGAAAGGAACG 19
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TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 89.5'
...ches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: H. sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-10-174-020-71/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-174-020-138
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                                                                      FEATURE:
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GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: McSwiggen, James

APPLICANT: McSwiggen, James

APPLICANT: Fosnaugh, Kathy

TITLE OF INVENTION: Gene Expression Using Short Interfering RNA

TITLE OF INVENTION: Gene Expression Using Short Interfering RNA

CURRENT APPLICATION NUMBER: US/10/224,005

CURRENT APPLICATION NUMBER: US 60/315,315

PRIOR APPLICATION NUMBER: US 60/315,315

PRIOR PILING DATE: 2001-08-28

SOFTWARE: PatentIn version 3.0

SEQ ID NO 315

LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region US-10-224-005-315
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Publication No. US2030170636A1
GENERAL INFORMATION:
TITLE OF INVENTION: ANTISENSE MODULATION OF JAGGED 2 EXPRESSION
TITLE REFERENCE: RTS-0244
CURRENT APPLICATION NUMBER: US/10/091,625
CURRENT FILING DATE: 2002-03-05
NUMBER OF SEQ ID NOS: 90
SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: KOLler, Erich
APPLICANT: Shepard, Peter J.
TITLE OF INVENTION: JAGGED 2 INHIBITORS FOR INDUCING APOPTOSIS
FILE REFERENCE: ISPH-0660
CURRENT APPLICATION NUMBER: US/10/096,399A
NUMBER OF SEQ ID NOS: 91
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 3.3%; Score 14.8; DB 1; Length 19; Best Local Similarity 88.9%; Pred. No. 2.2e+02; Matches 16; Conservative 0; Mismatches 2; Indels
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93.8%; Pred. No. 1.7e+02;
ative 0; Mismatches 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19 GGGAGGGAGGTGGCCGT 2
          US20030143732A1
                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      265 CCCGGGGCTTCTCCGG 280
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 cccacccrrcrcccc 16
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US-10-096-399A-4
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Best Local
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US-10-126-704-76

Sequence 76, Application US/10126704

Sequence 76, Application US/10126704

Publication No. US20030170647A1

GENERAL INFORMATION:

APPLICANT: Bougueleret, Lydie

TITLE OF INVENTION: A NUCLEIC ACID ENCODING A RETINOBLASTOMA BINDING PROTEIN (RBP-7)

TITLE OF INVENTION: AND POLYMORPHIC MARKERS ASSOCIATED WITH SAID NUCLEIC ACID.

FILE REFERENCE: 44.US5.DIV

CURRENT FILING DATE: 2002-04-20

PRIOR APPLICATION NUMBER: US 60/091,315

PRIOR FILING DATE: 1998-06-30

PRIOR FILING DATE: 1998-12-10

NUMBER OF SEQ ID NOS: 140

SOFTWARE: PATENT.

SEQ ID NO 76

LENGTH: 20

LENGTH: 20

LENGTH: 20
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APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: Fosnaugh, Kathy
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Adenosine Al Receptor (AD
TITLE OF INVENTION: Gene Expression Using Short Interfering RNA
TITLE REFERENCE: 900/414 (MBHB01-1110-A)
CURRENT APPLICATION NUMBER: US/10/224,005
RIOR APPLICATION NUMBER: US 60/315,315
PRIOR APPLICANT DATE: 2001-08-28
NUMBER OF SEQ ID NOS: 347
SOFTWARE: PatentIn version 3.0
SEQ ID NO 154
LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; LOCATION: 1..20
; OTHER INFORMATION: upstream amplification primer for SEQ 34, SEQ 55, SEQ 35, SEQ 56
US-10-126-704-76
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3.4%; Score 15.4; DB 1; Length 20;
Best Local Similarity 94.1%; Pred. No. 2.1e+02;
Matches 16; Conservative 0; Mismatches 1; Indels
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3.3%; Score 14.8; DB 1; Length 1
Best Local Similarity 77.8%; Pred. No. 2.2e+02;
Matches 14; Conservative 2; Mismatches 2; Indels
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Sequence 154, Application US/10224005
Publication No. US20030143732A1
GENERAL INFORMATION:
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US-10-224-005-315/c
; Sequence 315, Application US/10224005
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: primer_bind
LOCATION: 1..20
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JERURAL INFORMATION
MORELL APPLICANT: Cowsert, Lex M.
MAPLICANT: Monell, John
MAPLICANT: Monell, John
MAPLICANT: Sasmor, Henri M.
MAPLICANT: Brooks, Douglas G.
MAPLICANT: Brooks, Douglas G.
MAPLICANT: Watt, Jacqueline R.
MAPLICANT: Wortt, Jacqueline R.
MAPLICANT: Wortt, Jacqueline R.
MAPLICANT: Borchers, Alexander
MAPLICANT: Wortt, Jacqueline R.
MAPLICANT: Borchers, Alexander
MAPLICANT: MARCHION: MODULATION BY OLIGONUCLEOTIDES FOR GENE MODULATION
MITTLE OF INVENTION: GENERATION OF OLIGONUCLEOTIDES FOR GENE MODULATION
MITTLE OF INVENTION: GENERATION OF OLIGONUCLEOTIDES FOR GENE MODULATION
MITTLE OF INVENTION: GENERATION OF OLIGONUCLEOTIDES FOR GENE MODULATION
MITTLE OF SEQ ID NOS: 947
MUNBER OF SEQ ID NOS: 947
MARE: FESTESEQ for Windows Version 4.0
MARCHIOLICANT: 16
MARCHIOLICANT: 16
MARCHIOLICANT: MARCHIOLICANTE MARC
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| Publication No. US20030207839A1
| GENERAL INFORMATION:
| APPLICANT: SUBBAN | Prejet | APPLICANT: SUBAN | Prejet | TITLE OF INVENTION: ANTISENSE MODULATION OF JAGGED 2 EXPRESSION | FILE REFERENCE: RTS-024 | CURRENT APPLICATION NUMBER: US/10/461,668 | CURRENT APPLICATION NUMBER: US/10/091,625 | PRIOR APPLICATION NUMBER: US/10/091,625 | PRIOR APPLICATION NUMBER: US/10/091,625 | NUMBER OF SEQ ID NOS: 90 | SEQ ID NO 4 | LENGTH: 16
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llarity 93.8%; Pred. No. 1.7e+02;
Conservative 0; Mismatches 1; Indels
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3.2%; Score 14.4; DB 1;
Best Local Similarity 93.8%; Pred. No. 1.7e+02;
Matches 15; Conservative 0; Mismatches 1;
                                                                                                              OTHER INFORMATION: Synthetic PCR primer US-10-096-399A-4
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1 CCCAGGCTTCTCCGG 16
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TYPE: DNA
ORGANISM: Artificial Sequence
TYPE: DNA
ORGANISM: Artificial Sequence
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US-10-461-668-4
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Best Local Similarity
Matches 15; Conserva
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                                                                            FEATURE:
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APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Draper. Ken
APPLICANT: Draper. Ken
APPLICANT: Draper. Ken
APPLICANT: Draper. Ken
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relater
TITLE OF INVENTION: Levels of NP-Kappa B
TITLE OF INVENTION: Levels of NP-Kappa B
CURRENT APPLICATION NUMBER: US/09/864,785
CURRENT FILING DATE: 2001-05-23
NUMBER OF SEQ ID NOS: 3929
SOFTWARE: PatentIn version 3.0
SEQ ID NO 145
LENGTH: 17
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Sequence 146, Application US/09664785

Sequence 146, Application US/09664785

Patent No. US200201756841

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: Stinchcomb, Dan

APPLICANT: McSwiggen, Jim

TITLE OF INVENTION: Brymatic Nucleic Acid Treatment of Diseases or Conditions Relater

TITLE OF INVENTION: Brymatic Nucleic Acid Treatment of Diseases or Conditions Relater

TITLE OF INVENTION: Levels of NP-Kappa B

FILE REFERENCE: 400/022 (MBHB00-812-D)

CURRENT ETLING DATE: 2001-05-23

NUMBER OF SEQ ID NOS: 3929

SOFTWARE: Patentin version 3.0

SEQ ID NO 146

LENGTH: 17
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US-09-864-785-145
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                                                                                                 Query Match
3.2%; Score 14.4; DB 1; Length 16;
Best Local Similarity 93.8%; Pred. No. 1.7e+02;
Matches 15; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                   Sequence 145, Application US/09864785; Patent No. US20020177568A1
GENERAL INFORMATION:
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                                                                                                                                                                                                          265 CCCGGGGCTTCTCCGG 280
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ORGANISM: Artificial Sequence
FEATURE:
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ORGANISM: Artificial Seguence
FEATURE:
; FEATURE:
; OTHER INFORMATION: PCR Primer
US-10-388-263-377
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Best Local Similarity 75.0°
Matches 12; Conservative
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GENERAL INFORMATION

GENERAL INFORMATION

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: Sandberg, Jennifer

APPLICANT: Gardon, Gilad

APPLICANT: Gordon, Gilad

APPLICANT: Gordon, Gilad

APPLICANT: Stinchomb, Dames

APPLICANT: Stinchomb, Dames

APPLICANT: Stinchomb, Dames

APPLICANT: Stinchomb, Dames

APPLICANT: McSwiggen, James

APPLICANT: MCWERN: RECEPTOR FOR THE TREATMENT OF ANGIOGENESIS RELATED DISEASES AND (CURRENT TAILNG DATE: 1905-10-16

CURRENT APPLICATION NUMBER: US 60/005,974

PRIOR FILING DATE: 1995-08-10

PRIOR FILING DATE: 1995-08-10

PRIOR FILING DATE: 2000-11-10

PRIOR FILING DATE: 2001-05-29

PRIOR FILING DATE: 2001-05-29

PRIOR FILING DATE: 2001-05-29

PRIOR PLILOR DATE: 2001-05-29

PRIOR FILING DATE: 2001-05-29

PRIOR PLILOR DATE: 2001-05-29

PRIOR PLILOR DATE: 2001-05-29

PRIOR PLILOR DATE: 2001-05-29

PRIOR PLILOR DATE: 2001-05-05-03

NUMBER OF SEQ ID NOS: 5999

SEQ ID NO 491

LENGTH: 17
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Fublication No. US20030054365A1
Fublication No. US20030054365A1
Fublication No. US20030054365A1
Fublication No. US20030054365A1
Fublication No. US2003054365A1
FUBLICANT: Qiu, Gang
APPLICANT: Qiu, Gang
APPLICANT: Humphreys, Robert
TILE OF INVENTION: CANCER CELL VACCINE
FILE REFERENCE: U.S. Application 09/205,995, (CIE)
CURRENT FILING DATE: 2002-01-22
FRIOR APPLICATION NUMBER: 09/036,746
FRIOR APPLICATION NUMBER: 08/661,627
FRIOR FILING DATE: 1998-06-11
NUMBER OF SEQ ID NOS: 79
SOFTWARE: Patentin Ver: 2.0
SEQ ID NO 24
     US20040220128A1
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ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16 Greardeccarrirr 1
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Best Local Similarity 93.8
Matches 15, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: RNA
ORGANISM: Homo Sapiens
Publication No. US20 GENERAL INFORMATION:
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                                                                                                                                                                           Sequence 7502, Application US/10138674

Sequence 7502, Application US/10138674

GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Ribozywe Pharmaceuticals, Inc.
APPLICANT: Stanchcomb, Dan
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Method and Reagent for the Treatment of Diseases or Conditions Re
FILE REFERENCE: MHBB00-876-N (400/049)
CURRENT APPLICATION NUMBER: US/10/138,674
NUMBER OF SEQ ID NOS: 20822
SOFTWARE: Patentin version 3.0
LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 7502, Application US/10287949A

Sequence 7502, Application US/10287949A

Publication No. US20040102389A1

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: Ravco, Pam

APPLICANT: Barcolomb, Dan

APPLICANT: Stinchcomb, Dan

APPLICANT: Stinchcomb, Dan

APPLICANT: Levels of Vascular Endothelial Growth Factor Receptor

TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor

CURRENT APPLICATION NUMBER: US/10/287,949A

NUMBER OF SEQ ID NOS: 2083-04-11

SOFTWARE: PatentIn version 3.0

SEQ ID NO 7502
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3.2%; Score 14.4; DB 1; Length 17;
Best Local Similarity 93.8%; Pred. No. 1.9e+02;
Matches 15; Conservative 0; Mismatches 1; Indels
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US-10-712-633-491/c
; Sequence 491, Application US/10712633
     131 CCTCGGCCTGCCGCCT 146
                                    28 GIGGIGGCCATITITY 43
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; ORGANISM: Homo sapiens
US-10-138-674-7502
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ORGANISM: Homo sapiens
                                                                                                                                                              -10-138-674-7502/c
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  Length 17;
                                                                                                                                                                                                                                                                                                                                       APPLICANT: Zwick, Michael G.
Edington, Brent E.
McSwiggen, James A.
Merlo, Datricia Ann Owens
Guo, Lining
Skokut, Thomas A.
Young, Scott A.
Folkerts, Otto
Merlo, Donald J.
TITLE OF INVENTION: COMPOSITION AND METHODS FOR
MODULATION OF GENE EXPRESSION
IN PLANTS
NUMBER OF SEQUENCES: 1263
                                                   IndelB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   IndelB
Query Match
3.1%; Score 14; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 2.1e+02;
Matches 14; Conservative 0; Mismatches 0;
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3.1%; Score 13.8; DB 1;
Best Local Similarity 64.7%; Pred. No. 2.2e+02;
Matches 11; Conservative 4; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P. C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/961,077
FILING DATE: 31-Sep-2001
CLASSIFICATION: «Unknown.»
PRIOR APPLICATION NUMBER: 08/679,645
FILING DATE: July 12, 1996
APPLICATION NUMBER: 60/001,135
FILING DATE: July 13, 1995
APPLICATION NUMBER: 60/001,26
FILING DATE: September 2, 1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 219/247
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE DESCRIPTION: SEQ ID NO: 826
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Lyon & Lyon
STREET: 633 West Pifth Street
Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Warburg, Richard J. REGISTRATION NUMBER: 32,327
                                                                                                                                                                                                                                           US-09-961-077-826; Sequence 826, Application US/09961077; Sequence 826, Application No. US20030014775A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 826:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS
                                                                                                   328 CTCTCGGGGGCGAG 341
                                                                                                                                               2 crcrceeeeccac 15
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                                                                                                                                                                                                                                   Sequence 2.2. Application US/10093958

Publication No. US2003004442341

GENERAL INFORMATION:
APPLICANT: Gillies, Stephen
APPLICANT: Gillies, Stephen
TITLE OF INVENTION: Expression Technology for Proteins Containing a Hybrid Isotype Ar
TITLE OF INVENTION: Expression Technology for Proteins Containing a Hybrid Isotype Ar
TITLE OF INVENTION: UMBER: US/10/093,958
CURRENT APPLICATION NUMBER: US/10/093,958
CURRENT FILING DATE: 2002-03-07
PRIOR FILING DATE: 2001-03-07
NUMBER OF SEQ ID NOS: 50
SOFTWARE: Patentin version 3.0
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APPLICANT: Maher, Majella
APPLICANT: Maher, Majella
APPLICANT: Martin, Cara
APPLICANT: Jannes, Geart
APPLICANT: Van Dor Weide, Marjo
APPLICANT: Van Dor Weide, Marjo
APPLICANT: Van Dor Weide, Marjo
TITLE OF INVENTION: Clinically important fungal pathogens
FILING DATE: 2004-10-19
PRIOR PILING DATE: 2000-05-24
PRIOR PILING DATE: 1999-06-28
PRIOR PILING DATE: 1999-06-28
PRIOR PILING DATE: 1999-06-11
NUMBER OF SEQ ID NOS: 43
SOFTWARE: PatentIn Ver: 2.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; OTHER INFORMATION: forward primer for gamma 1 hinge region US-10-093-958-22
                            1; Indels
  93.8%; Pred. No. 2.1e+02;
tive 0; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; Sequence 21, Application US/10967254; Publication No. US20050164243A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; TYPE: DNA
; ORGANISM: Aspergillus versicolor
US-10-967-254-21
                                                                            220 GGTCGCCTGCCCAGCC 235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               300 GAAGAGTTGGGCTCTG 315
                                                                                                        1 GGTGGCCTGCCCAGCC 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA ORGANISM: artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19 GAAGATTTGGGCTCTG 4
Best Local Similarity 93.8 Matches 15; Conservative
                                                                                                                                                                                                                         US-10-093-958-22/c
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RESULT 251
US-09-930-423-1471
US-09-930-423-1471
Sequence 1471, Application US/09930423
Fublication No. US2003009203A1
FUBLICANT: Ribozywe Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: MCSWiggen, Jim
FILE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease
FILE REPERENCE: MBHB00,918-A 400/027
CURRENT PELING DATE: 2001-08-15
NUMBER OF SEQ ID NOS: 4553
SOFTWARE PARENTIN Version 3.0
SEQ ID NO 1471
LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 1477. Application US/09930423
Fublication No. US20030092003A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease
FILE REFERENCE: WAHBOO, 918-A 400,027
CURRENT APPLICATION NUMBER: US/09/930,423
CURRENT FILNG DATE: 2001-08-15
SOFTWARR: Patentin version 3.0
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SOFTWARR: Patentin version 3.0
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                                                                                                                                                                                                                          3.1%; Score 13.8; DB 1; Length 17; 88.2%; Pred. No. 2.2e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 3.1%; Score 13.8; DB 1; Length 17; Best Local Similarity 88.2%; Pred. No. 2.2e+02; Matches 15; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                         Indels
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3.1%; Score 13.8; DB 1;
Best Local Similarity 88.2%; Pred. No. 2.2e+02;
Matches 15; Conservative 0; Mismatches 2;
            2001-08-15
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       CURRENT FILING DATE: 2001-08-1
NUMBER OF SEQ ID NOS: 4553
SOFTWARE: PatentIn version 3.0
SEQ ID NO 336
LENGTH: 17
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Matches 15, Conservative
                                                                                                                           ; TYPE: RNA
; ORGANISM: Homo Sapiens
US-09-930-423-336
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US-09-930-423-1471
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US-09-930-423-1472
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US-09-930-423-1472
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                                                                                                          RESULT 248
US-09-848-754A-3581
US-09-848-754A-3581
US-09-848-754A-3581

SABIL SABIL APPLICATION US/09848754A

PUBLICATION NO. US20030073207A1

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc.

TITLE OF INVENTION: Levels of Epidermal Growth Factor Receptors

TITLE OF INVENTION: Levels of Epidermal Growth Factor Receptors

CURRENT APPLICATION NUMBER: US/09/848,754A

CURRENT FILING DATE: 2001-05-03

NUMBER OF SEQ ID NOS: 9645

SOFTWARE: PatentIn version 3.0

SEQ ID NO 3581
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Publication No. US200300920031
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blat, Larry
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease
FILE REFERENCE: MaHBO0, 918-A 400/027
CURRENT PILING DATE: 2001-08-15
NUMBER OF SEQ ID NOS: 4553
SOFTWARE: Patentin version 3.0
SEQ ID NO 5
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Publication No. US20030092003A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Matt, Larry
APPLICANT: Makingan, Jim
TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease
FILE REFERENCE: MBHB00,918-A 400/027
CURRENT APPLICATION NUMBER: US/09/930,423
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3.1%; Score 13.8; DB 1;
Best Local Similarity 88.2%; Pred. No. 2.2e+02;
Matches 15; Conservative 0; Mismatches 2;
106 CGCTGACTTTCAGCGGG 122
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                           1 CGCUGCCUUUCAGCUGG 17
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Matches 15, Conservative
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ORGANISM: Homo sapiens
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ORGANISM: Homo Sapiens
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US-09-930-423-336/c
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APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relater
TITLE OF INVENTION: Hepatitis C Virus Infection
FITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: RPI 400/09/740,332
CURRENT APPLICATION NUMBER: US/09/740,332
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9704
SOFTWARE: Patentin version 3.0
SEQ ID NO 3130
LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 129, Application US/09792818

Sequence 129, Application US/09792818

Sequence 129, Application No. US20030134806A1

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: Jarvis, Thale

APPLICANT: McSwiggen, Jim

APPLICANT: McSwiggen, Jim

APPLICANT: McSwiggen, Jim

APPLICANT: McSwiggen, Jim

APPLICANT: Bilis, Jonathan

TITLE OF INVENTION: McShod and Reagent for the Inhibition of Grb-2-related with Insertication NUMBER: US/09/792,818

CURRENT APPLICATION NUMBER: US/09/792,818

CURRENT APPLICATION NUMBER: US/09/792,818

SOFTWARE: Patentin version 3.0

SOFTWARE: Patentin version 3.0

TITLE OF INVENTION: 2304

SOFTWARE: Patentin version 3.0
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APPLICANT: Von Carlowitz, Ira
APPLICANT: McSwiggen, Jim
APPLICANT: Hamblin, Paul
APPLICANT: Ellis, Jonathan
APPLICANT: Ellis, Jonathan
TITLE OF INVENTION: Method and Reagent for the Inhibition of Grb-2-related with Inser
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88.2%; Pred. No. 2.2e+02;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: oligonucleotide substrate
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 330, Application US/09792818 Publication No. US20030134806A1 GENERAL INFORMATION:
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Best Local Similarity 88.4.
These 15; Conservative
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Best Local Similarity 88.29
Matches 15; Conservative
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NAME/KEY: misc_feature
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ORGANISM: Homo sapiens
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US-09-792-818-330/c
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                                                                          US-09-827-395A-630

i Sequence 630, Application US/09827395A

i Sequence 630, Application US/09827395A

i Publication No. US20030113891A1

i GENERAL INFORMATION:

APPLICANT: Barare Chownina

APPLICANT: Daws MCSWiggen

APPLICANT: Bharar Chownina

ITILE OF INVENTION: Method and Reagent for the Inhibition of NOGO Receptor G

FILE REFRENCE: MBHB00-99-80-C (400/017)

CURRENT APPLICATION NUMBER: US/09/027,395A

CURRENT APPLICATION NUMBER: US/09/027,395A

CURRENT PILING DATE: 2001-04-05

PRIOR APPLICATION NUMBER: 60/181,797

PRIOR PILING DATE: 2000-02-11

NUMBER OF SEQ ID NOS: 2617

SOFTWARE: PatentIn version 3.0

SEQ ID NO 630

I DN O 630
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Publication No. US20030125270A1

GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: Hepatitis C Virus Infection
TITLE OF INVENTION: Hepatitis C Virus Infection
TITLE REPERENCE: RP1 400/003
CURRENT APPLICATION NUMBER: US/09/740,332
CURRENT FILING DATE: 2001-03-26

NUMBER OF SEQ ID NOS: 9704
SEQ ID NO 800
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  1 CCCGCCGGGAGCCCGCG 17
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ORGANISM: artificial sequence
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ORGANISM: Homo sapiens
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Matches 13; Conserv
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US-09-740-332-3130/c
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242 CCCCGCCTGGAGGCCGC 258
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Best Local Similarity 88.2
Matches 15; Conservative
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US-09-745-237A-1471
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ORGANISM: Homo sapiens
                                                                                                                                              RESULT 260
US-09-745-237A-1471
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Bublication No. US20030143708A1

GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: Missurgen, Jim
TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease
FILE REFERENCE: 400/007 (MBHB00-918-A)
CURRENT FILING DATE: 2002-04-15
NUMBER OF SEQ ID NOS: 4550
SEQ ID NO 5
SEQ ID NO 5
LENGTH: 17
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PUBLICATION No. US20030143708A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: Machine Method and Reagent for the Treatment of Alzheimer's Disease;
TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease;
CURRENT APPLICATION NUMBER: US/09/745, 237A
NUMBER OF SEQ ID NOS: 4550
SOFTWARE PATELING DATE: 2002-04-15
NUMBER OF SEQ ID NOS: 4550
SOFTWARE PATENTING PATE: 2002-04-15
SOFTWARE PATE: 2002-04-15
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88.2%; Pred. No. 2.2e+02;
Live 0; Mismatches 2; Indels
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3.1%; Score 13.8; DB 1; Length 17;
Best Local Similarity 88.2%; Pred. No. 2.2e+02;
Matches 15; Conservative 0; Mismatches 2; Indels
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88.2%; Pred. No. 2.2e+02;
tive 0; Mismatches 2;
                FILE REFERENCE: MHHB00-901-A (400/013)
CURRENT APPLICATION NUMBER: US/09/792,818
CURRENT FILING DATE: 2001-02-23
NUMBER OF SEQ ID NOS: 2304
SOFTWARE: PatentIn version 3.0
TITLE OF INVENTION: (GRID) Gene
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                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 88.2
Matches 15; Conservative
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Matches 15; Conservative
                                                                                                                                                                   ; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-792-818-330
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ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
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LENGTH: 17
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US-05-817-879-800
US-05-817-879-800
Sequence 800, Application US/09817879
FUBLICATION NO. US20030171311A1
GENERAL INFORMATION:
APPLICANT: Ribosyme Pharmaceuticals Inc.
TITLE OF INVENTION: ADDITION: ADDITIONS Related TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: MBHB00-801-F
Sequence 1471, Application US/09745237A

Sequence 1471, Application US/09745237A

Publication No. US20030143708A1

SEMERAL INFORMATION:

APPLICANT: Ribory Pharmaceuticals, Inc.

APPLICANT: Blatt, Larry

APPLICANT: McSwiggen, Jim

TITLE OF INVENTION: McHabod and Reagent for the Treatment of Alzheimer's Disease

CURRENT FILMS DATE: 2002-04-15

CURRENT FILMS DATE: 2002-04-15

NUMBER OF SEQ ID NOS: 4550

SOFTWARE: Patentin version 3.0

SOFTWARE: Patentin version 3.0

SOFTWARE: The Company of the Comp
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Publication No. US20030143708A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blact, Larry
APPLICANT: Blact, Larry
APPLICANT: Machod and Reagent for the Treatment of Alzheimer's Disease
UURENT APPLICATION UMBER: US/09/745,237A
CURRENT FILING DATE: 2002-04-15
NUMBER OF SEQ ID NOS: 4550
SSOTUMARE PATORICATION TOWN OF TANGE SEQ ID NO 1450
LENGTH: 17
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Sequence 5872, Application US/10156306

Publication No. US20030119017A1

Publication No. US20030119017A1

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: McSwiggen, James

TITLE OF INVENTION: Levels of IKK-Gamma and PKR

FILE REPERENCE: MBHB01-664-A (400/050)

CURRENT APPLICATION NUMBER: US/10/156,306

CURRENT FILING DATE: 2002-05-28

NUMBER OF SEQ ID NOS: 8013

SOFTWARE: Patentin version 3.0

SEQ ID NO 5872
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PUBLICANT: Application No. US20030191077A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Rosnaugh, Kathy
APPLICANT: Rosnaugh, Jim
TITLE OF INVENTION: METHOD AND REAGENT FOR THE TREATMENT OF ASTHMA AND ALLERGIC CONDI
FILE REPERRORS: 400/056 (MBHB01-1110)
FILE REPERRORS: 100/056 (MBHB01-1110)
FILE OF INVENTION NUMBER: US 60/315,315
PRIOR APPLICATION NUMBER: US 60/315,315
PRIOR APPLICATION NUMBER: US 60/315,315
SOFTWARE: PARCELLIN VERSION 3.0
SOFTWARE: PARCELLIN VERSION 3.0
SSOID NO. 2217
TENNING 12177
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                                             Query Match
3.1%; Score 13.8; DB 1;
Best Local Similarity 76.5%; Pred. No. 2.2e+02;
Matches 13; Conservative 2; Mismatches 2;
                                                                                                                                                 246 GCCTGGAGGCCGCGGTC 262
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Best Local Similarity 76.55
These 13; Conservative
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Best Local Similarity 88.2
Matches 15; Conservative
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; ORGANISM: Homo sapiens
US-10-156-306-5872
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CORGANISM: Homo sapiens
US-10-230-006-2217
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US-10-230-006-2217/c
     US-10-156-306-5871
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US-10-156-306-5872
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| Publication No. US20030119017A1 |
| GENERAL INFORMATION: |
| APPLICANT: Ribozyme Pharmaceuticals, Inc. |
| APPLICANT: Ribozyme Pharmaceuticals, Inc. |
| APPLICANT: MCSWiggen, James |
| TITLE OF INVENTION: Levels of IKK-Gamma and PKR |
| TITLE OF INVENTION: Levels of IKK-Gamma and PKR |
| TITLE OF INVENTION: Levels of IKK-Gamma and PKR |
| TITLE OF INVENTION: Levels of IKK-Gamma and PKR |
| TITLE OF INVENTION UNMBER: US/10/156,306 |
| CURRENT PLING DATE: 2002-05-28 |
| NUMBER OF SEQ ID NOS: 8013 |
| SOFTWARE: PatentIn version 3.0 |
| SEQ ID NO 5871 |
| LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: MBH800-801-F
CURRENT APPLICATION WUMBER: US/09/817,879
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9703
SOFTWARE: Patentin version 3.0
SEQ ID NO 3130
LENGTH: 17
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                                                                                                                                                                                          FEATURE:
NAME/KEY: misc_feature
LOCATION:
PEATURE:
LOCATION:
OTHER INFORMATION: oligonucleotide substrate
US-09-817-879-800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: oligonucleotide substrate
CURRENT APPLICATION NUMBER: US/09/817,879
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9703
SOFTWARE: PatentIn version 3.0
LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Sequence 3130, Application US/09817879; Publication No. US20030171311A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                             432 AGGACTCGGCTCACACA 448
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ORGANISM: artificial sequence
                                                                                                                                                 TYPE: RNA
ORGANISM: artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
NAME/KEY: misc_feature
LOCATION:
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-817-879-3130/c
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US-10-156-306-5871
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APPLICANT: Pamela, Pavco,
APPLICANT: Pamela, Pavco
APPLICANT: Pamela, Pavco
APPLICANT: Pamela, Pavco
APPLICANT: Patrice, Lee
APPLICANT: Patrice, Lee
APPLICANT: Elsabeth, Daper
APPLICANT: Elsabeth, Daper
TITLE OF INVENTION: OLGONOLICEOTIDE MEDIATED INHIBITION OF HEPATITIS B VIRUS AND HEPAN
TITLE OF INVENTION: OLGONOLICEOTIDE MEDIATED INHIBITION OF HEPATITIS B VIRUS AND HEPAN
TITLE OF INVENTION: OLGONOLICEOTIDE MEDIATED INHIBITION OF HEPATITIS B VIRUS APPLICATION NUMBER: US /10/669, 841
CURRENT PELING DATE: 2003-09-23
PRIOR PELING DAPE: 2003-09-26
PRIOR PALING DAPE: 2001-06-66
PRIOR APPLICATION NUMBER: US 60/335, 059
PRIOR PILING DAPE: 2001-10-66
PRIOR PELING DAPE: 2001-10-67
PRIOR PELING DAPE: 2002-03-10
PRIOR PILING DAPE: 2002-03-11
PRIOR PILING DAPE: 2002-03-11
PRIOR PILING DAPE: 2002-03-11
PRIOR PILING DAPE: 2000-07-07
PRIOR PELING DAPE: 2000-07-07
PRIOR APPLICATION NUMBER: US 69/504,321
PRIOR PILING DAPE: 2000-07-07
PRIOR APPLICATION NUMBER: US 69/504,321
PRIOR PILING DAPE: 2000-07-07
PRIOR PILING DAPE: 2000-07-07
PRIOR PRIOR PILING DAPE: 2000-07-07
PRIOR PILING DAPE: 2000-07-07
PRIOR PILING DAPE: 2000-07-07
PRIOR PILING DAPE: 2000-07-07
PRIOR PILING DAPE: 2000-02-15
PRIOR PILING DAPE: 2000-07-07
PRIOR PILING DAPE: PADELICAND DAPE: PADELICA
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                                                                                                                                                         Sequence 3393, Application US/10669841 Publication No. US20040127446A1 GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc. APPLICANT: Lawrence, Blatt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-10-669-841-5723/C
Sequence 5723, Application US/10669841
Sequence 5723, Application US/10669841
GENERAL INFORMATION:
APPLICANT: STATE Therespectics, Inc.
APPLICANT: Lawrence, Blatt
APPLICANT: Dennis, Macejak
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                          Dennis, Macejak
James, McSwiggen
David, Morrissey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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Best Local Similarity
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Matches
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                                                                         GENERAL INFORMATION:

APPLICANT: Hibozyme Pharmaceuticals, Inc.
APPLICANT: Hibozyme Blatt
APPLICANT: Gawrence Blatt
APPLICANT: Bararat Chowrira
APPLICANT: Bharat Chowrira
CURRENT PILING DATE: 2003-05-06
FILE REPERENCE: MBHB00-3F8-H (400/112)
CURRENT FILING DATE: 2001-04-05
FRIOR APPLICATION NUMBER: 09/827,395
FRIOR PILING DATE: 2001-02-09
FRIOR PILING DATE: 2001-02-09
FRIOR PILING DATE: 2000-02-01
FRIOR APPLICATION NUMBER: 60/181,797
FRIOR FILING DATE: 2000-02-01
FRIOR FILING DATE: 2000-02-01
FRIOR FILING DATE: 2000-02-03
NUMBER OF SEQ ID NOS: 2617
SOFTWARE: PatentIn version 3.0
FRIOR FILING DATE: 2002-04-03
NUMBER OF SEQ ID NOS: 2617
SEQ ID NO 630
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 2726, Application US/10712672

Sequence 2726, Application US/10712672

Publication No. US20040102413A1

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharaceuticals, Inc.

APPLICANT: Chowriza, Bharat

APPLICANT: Chowriza, Bharat

APPLICANT: Chowriza, Bharat

APPLICANT: Stinchcomb, Dan

TITLE OF INVENTION: Method and Reagent for the Inhibition of Telomerase Enzyme

FILE REFERENCE: MBHB00-082-C (400/019)

CURRENT FILING DATE: 2003-11-13

FRIOR APPLICATION NUMBER: US/10/12,672

CURRENT FILING DATE: 2000-08-31

PRIOR APPLICATION NUMBER: 60/197,769

PRIOR PILING DATE: 2000-04-14

PRIOR PILING DATE: 1999-08-31

PRIOR FILING DATE: 1999-08-31

NUMBER OF SEQ ID NOS: 5586

SOFTWARE: Patentin version 3.0

FROGUL DO 2726
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3.1%; Score 13.8; DB 1; Length 17;
Best Local Similarity 88.2%; Pred. No. 2.2e+02;
Matches 15; Conservative 0; Mismatches 2; Indels
Sequence 630, Application US/10430882
Publication No. US20030203870A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         263 GGCCCGGGCCTTCTCCG 279
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 GGCCCGGGCCCGUCCG 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-430-882-630
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-712-672-2726
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US-10-712-672-2726/c
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Sequence 27, Application US/10158160A
; Bublication No. US20030059805A1
; GENERAL INFORMATION:
    GENERAL INFORMATION:
    APPLICANT: RAPPOLD-HOERBRAND, GUDRUN
    APPLICANT: RAPPOLD-HOERBRAND, GUDRUN
    APPLICANT: RAPPOLD-HOERBRAND, GUDRUN
    TITLE OF INVENTION: HUMAN GROWTH GENE AND SHORT STATURE GENE REGION
    TITLE OF INVENTION: HUMAN GROWTH GENE AND SHORT STATURE GENE REGION
    TITLE OF INVENTION: HUMAN GROWTH GENE AND SHORT STATURE GENE REGION
    TILLE OF INVENTION: HUMANER: 09/147,699
    CURRENT FILING DATE: 1099-06-24
    PRIOR APPLICATION NUMBER: 00/27,633
    PRIOR PELING DATE: 1997-09-29
    PRIOR PELING DATE: 1996-10-01
    PRIOR PILING DATE: 1996-10-01
    PRIOR PELING DATE: 1996-10-01
    PRIOR PILING DATE: 1997-01-16
    NUMBER OF SEQ ID NOS: 55
    SEQ ID NO 27
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                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: WORD Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/961,077
FILING DATE: 21-890-2001
CLASSIFICATION: «Unknown»
PRIOR APPLICATION NUMBER: 08/69,645
FILING DATE: July 12, 1996
APPLICATION NUMBER: 60/001,135
FILING DATE: July 13, 1995
APPLICATION NUMBER: 09/001,135
FILING DATE: SEPTEMBER: 09/001,135
FILING DATE: SEPTEMBER: 09/001,135
FILING DATE: SEPTEMBER: 09/001,135
FILING DATE: SEPTEMBER: 09/001,135
                                                                                                                                                                                                   COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 219/247
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear sequence DESCRIPTION: SEQ ID NO: 571: US-09-961-077-571
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: 643 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 571:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                        CITY: Los Angeles
STATE: California
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Matches 14; Conservat
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                                                                                                               APPLICANT: Renneth, Draper
APPLICANT: Elisabeth, Roberte
TITLE OF INVENTION OLIGOMOCLEOTIDE MEDIATED INHIBITION OF HEPATITIS B VIRUS AND HEPA
TITLE OF INVENTION: OLIGOMOCLEOTIDE MEDIATED INHIBITION OF HEPATITIS B VIRUS AND HEPA
TITLE OF INVENTION: VIRUS REPLICATION
FILE REFERENCE: 400/042US (WHHB02-249-E)
CURRENT APPLICATION NUMBER: US/10/669,841
CURRENT APPLICATION NUMBER: US/2003-09-26
PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-00-24
PRIOR PELICATION NUMBER: US 60/335,059
PRIOR FILING DATE: 2001-10-24
PRIOR FILING DATE: 2001-10-24
PRIOR FILING DATE: 2001-12-05
PRIOR FILING DATE: 2001-12-06
PRIOR FILING DATE: 2001-12-06
PRIOR FILING DATE: 2001-03-11
PRIOR FILING DATE: 2001-03-26
PRIOR FILING DATE: 2001-03-26
PRIOR FILING DATE: 2001-03-26
PRIOR FILING DATE: 2000-03-11
PRIOR FILING DATE: 2000-03-11
PRIOR FILICATION NUMBER: US 09/740,332
PRIOR FILING DATE: 2000-03-16
PRIOR FILING DATE: 2000-03-15
PRIOR PRIOR
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OTHER INFORMATION: Description of Artificial Sequence: Nucleic Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
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NUMBER OF SEQUENCES: 1263
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 511, Application US/09961077
Publication No. US20030014775A1
GENERAL INFORMATION:
APPLICANT: Edington, Brent E.
MCSWiggen, James A.
MCSWiggen, James A.
Merlo, Eatricia Ann Owens
Guo, Lining
Skokut, Thomas A.
Young, Scott A.
FOLKerts, Otto
Merlo, Donald J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ), OTHER INFORMATION: oligonucleotide substrate US-10-669-841-5723
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: RNA
ORGANISM: Artificial Sequence
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                                                                            Pamela, Pavco
                                                                                                        Patrice, Lee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: misc_feature
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US-09-961-077-571
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US-09-740-332-3638/c

Sequence 3638, Application US/09740332

Sequence 3638, Application World US/09740332

Publication No. US203012527041

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals Inc.

TITLE OF INVENTION: Hepatitis C Virus Infection

FILE REPERENCE: RPI 400/003

CURRENT FILING DATE: 2001-03-26

NUMBER OF SEQ ID NOS: 9704

SOFTWARE: Patentin Version 3.0

TEMBER OF SEQ ID NOS: 9704

SOFTWARE: PATENTIN VERSION 3.0

SEQ ID NO 3638
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 276

US-09-792-818-429/C

Sequence 429, Application US/09792818

Sequence 429, Application No. US20030134806A1

Sequence 429, Application No. US20030134806A1

SEQUENCE 429, Application No. US20030134806A1

SEPLICANT: North Carlowitz, Ira

APPLICANT: Use Carlowitz, Ira

APPLICANT: Workswigen, Jim

APPLICANT: Hamblin, Paul

TITLE OF INVENTION: Method and Reagent for the Inhibition of Grb-2-related with Insert

TITLE OF INVENTION: (GRID) Gene

TITLE OF INVENTION: (GRID) Gene

TITLE OF INVENTION: (AND GRID) Gene

TITLE OF INVENTION: (AND GRID) Gene

CURRENT FILING DATE: 2001-02-23

NUMBER OF SEQ ID NOS: 2304

SEQ ID NO 429

LENGTH: 17
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                      3.0%; Score 13.4; DB 1; Length 17; 80.0%; Pred. No. 2.4e+02; ive 2; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
3.0%; Score 13.4; DB 1; Length 17;
Best Local Similarity 93.3%; Pred. No. 2.4e+02;
Matches 14; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 17;
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3.0%; Score 13.4; DB 1;
Best Local Similarity 93.3%; Pred. No. 2.4e+02;
Matches 14; Conservative 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; OTHER INFORMATION: oligonucleotide substrate US-09-740-332-3638
                                                                                                               210 ACCTGCGGCGGGTCG 224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: RNA
ORGANISM: artificial sequence
                                                                                                                                             2 ACCUGCGCCGCCCC 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       210 ACCTGCGGCGGGTCG 224
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                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: RNA
ORGANISM: Homo sapiens
                 Query Match
Best Local Similarity
Matches 12, Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-792-818-429
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Sequence 1488, Application US/09864785

Sequence 1488, Application US/09864785

Sequence 1488, Application US/09864785

GENERAL INFORMATION:
APPLICANT: Stinchcomb, Dan
APPLICANT: Draper, Ken
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Levels of NF-Kappa B
FILE REFERENCE 140/022 (MBH000-812-D)
CURRENT APPLICATION UNMERS: US/09/864,785

CURRENT APPLICATION UNMERS: 2001-05-23

NUMBER OF SEQ ID NOS: 3929

SOFTWARE PATENTING DATE: 2001-05-23

NUMBER OF SEQ ID NOS: 3329
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals Inc.
APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Bazymetic Nucleic Acid Treatment of Diseases or Conditions Relate TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: RPI 400/003
CURRENT APPLICATION WUMBER: US/09/740,332
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9704
SOFTWARE: Patentin version 3.0
LENGTH: 17
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US-09-864-785-1488
                                                                                         OTHER INFORMATION: Description of Artificial Sequence: synthetic CTHER INFORMATION: primer US-10-158-160A-27
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                                                                                                                                                                                            3.1%; Score 13.8; DB 1; Length 18; 88.2%; Pred. No. 2.4e+02; tive 0; Mismatches 2; Indels
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3.0%; Score 13.4; DB 1; Length 17;
Best Local Similarity 73.3%; Pred. No. 2.4e+02;
Matches 11; Conservative 3; Mismatches 1; Indels
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. OTHER INFORMATION: oligonucleotide substrate
US-09-740-332-917
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US-09-740-332-917
Sequence 917, Application US/09740332
Publication No. US20030125270A1
                                                                                                                                                                                                                                                                                      332 CGGGGGGGGGGGGAGG 348
                        TYPE: DNA ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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ORGANISM: artificial sequence
FEATURE:
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Best Local Similarity 88.2°
Matches 15; Conservative
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LENGTH: 17
LENGTH: 18
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US-10-232-634-12

Sequence 12, Application US/10232634

Publication No. US20030105314A1

GENERAL INFORMATION:

APPLICANT: Guida, Marco

APPLICANT: Hall, Jeff

TITLE OF INVENTION: GENERIC TYPING OF THE HUMAN CYTOCHROME P450 2A6 GENE

TITLE OF INVENTION: AND RELATED MATERIALS AND METHODS

FILE REFERENCE: 4389-20

CURRENT APPLICATION NUMBER: US/10/232,634

CURRENT PLIING DATE: 2002-08-30

PRIOR APPLICATION NUMBER: US/09/586,376

PRIOR RELING DATE: 2000-06-02

NUMBER OF SEQ ID NOS: 29

SEQ ID NO 12
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| Publication No. US20030105314A1
| GENERAL INFORMATION |
| APPLICANT: Gaida, Marco |
| APPLICANT: Gaida, Marco |
| APPLICANT: Hall, Jeff |
| TITLE OF INVENTION: GENETIC TYPING OF THE HUMAN CYTOCHROME P450 2A6 GENE |
| TITLE OF INVENTION: AND RELATED MATERIALS AND METHODS |
| FILE REFERENCE: 4389-20 |
| CURRENT APPLICATION NUMBER: US/10/232,634 |
| CURRENT PILING DATE: 2002-08-30 |
| PRIOR PILICATION NUMBER: US/09/586,376 |
| NUMBER OF SEQ ID NOS: 29 |
| SOFTWARE: Patentin Ver. 2.1
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3.0%; Score 13.4; DB 1;
Best Local Similarity 93.3%; Pred. No. 2.4e+02;
Matches 14; Conservative 0; Mismatches 1;
TITLE OF INVENTION: Hepatitis C Virus Infection FILE REFERENCE: MBHB00-801-F CURRENT APPLICATION NUMBER: US/09/817,879 CURRENT FILING DATE: 2001-03-26 NUMBER OF SEQ ID NOS: 9703 SSCTWARE: Patentin version 3.0 SEQ ID NO 3638 LENGTH: 17
                                                                                                                                                                                                                                                                                                                     ; OTHER INFORMATION: oligonucleotide substrate US-09-817-879-3638
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ORGANISM: Homo sapiens
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Best Local Similarity
Matches 14; Conserv
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US-10-232-634-11
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                                                                                                                                                                                                         TYPE: RNA
                                                                                                                                                                                                                                                          FEATURE
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                                                                                                                                                               APPLICANT: Jarvis, Thaie
APPLICANT: Von Carlowitz, Ira
APPLICANT: Won Carlowitz, Ira
APPLICANT: Won Carlowitz, Ira
APPLICANT: Won Carlowitz, Jim
APPLICANT: Ellis, Jonathan
TITLE OF INVENTION: Method and Reagent for the Inhibition of Grb-2-related with Inser
TITLE OF INVENTION: (GRID) Gene
FILE REFERENCE: MBHB00-901-A (400/013)
CURRENT APPLICATION NUMBER: US/09/792,818
CURRENT APPLICATION NUMBER: US/09/792,818
NUMBER OF SEQ ID NOS: 2304
SOFTWARE: PatentIn version 3.0
SOFTWARE: PatentIn version 3.0
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Publication No. US20030171311A1
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: MBHB00-801-F
CURRENT APPLICATION WMBER: US/09/817,879
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9703
SOFTWARE: PatentIn version 3.0
SEQ ID NO 917
LENGTH: 17
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; Publication No. US20030171311A1
; GENERAL INFORMATION:
; APPLICANT: Ribozyme Pharmaceuticals Inc.
; TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
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3.0%; Score 13.4; DB 1;
Best Local Similarity 93.3%; Pred. No. 2.4e+02;
Matches 14; Conservative 0; Mismatches 1;
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                                                                        Sequence 857, Application US/09792818
Publication No. US20030134806A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                132 CTCGGCCTGCCGCT 146
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CAGANISM: Homo sapiens
US-09-792-818-857
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                                                 US-09-792-818-857/c
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US-10-157-580A-68
Sequence 68, Application US/10157580A
Publication No. US20030124513A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, Jim.
TITLE OF INVENTION: Related To Levels Of HIV
TITLE OF INVENTION: Related To Levels Of HIV
TITLE OF INVENTION: Related To Levels Of HIV
CURRENT APPLICATION WUMBER: US/10/157,580A
CURRENT APPLICATION WUMBER: US/10/157,580A
NUMBER OF SEQ ID NOS: 170
SOFTWARE: Patentin version 3.0
SOFTWARE: Patentin version 3.0
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US-10-138-674-4559

Sequence 4559, Application US/10138674

Publication No. US20040077565A1

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: Racobed, Jaime

APPLICANT: Escobed, Jaime

TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Rell

TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor

CURRENT APPLICATION NUMBER: US/10/138,674

CURRENT FILING DATE: 2002-05-03

NUMBER OF SEQ ID NOS: 20822

SOFTWARE: Patentin version 3.0

TENDER OF SEQ ID NOS: 20822

SEQ ID NO 4559
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Publication No. US20040077565A1

GENERAL INFORMATION:
APPLICANT: Bavco, Pam
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stochedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Rel
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00-876-N (400/049)
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3.0%; Score 13.4; DB 1; Length 17;
Best Local Similarity 80.0%; Pred. No. 2.4e+02;
Matches 12; Conservative 2; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                       ; ORGANISM: Human immunodeficiency virus US-10-157-580A-68
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; ORGANISM: Homo sapiens
US-10-138-674-4559
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Best Local Similarity
Matches 13; Conserv
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US-10-157-580A-38
Sequence 38, Application US/10157580A
Sequence 38, Application US/10157580A
Publication No. US20030124513A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: RESuggen, Jim
ITITLE OF INVENTION: Related To Levels Of HIV
FILLE REPRENCE: MBH801-665-A (400/051)
CURRENT APPLICATION NUMBER: US/10/157,580A
CURRENT FILING DATE: 2002-08-30
NUMBER OF SEQ ID NOS: 170
SOFTWARE: Patentin version 3.0
SEQ ID NO 38
FRNGTH: 17
                                                                                                                                                                                                                                                                                                                                                                              Sequence 27, Application US/10157580A
Sequence 27, Application US/10157580A
Publication No. US20030124513A1
GENERAL INFORMATION:
BEPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Risymatic Nucleic Acid Treatment of Diseases of Diseases or CondiTITLE OF INVENTION: Related To Levels of HIV
FILE OF INVENTION: Related To Levels of HIV
FILE REPERENCE: MBHB01-66-A (400/051)
CURRENT APPLICATION NUMBER: US/10/157,580A
CURRENT FILING DATE: 2002-08-30
SOFTWARE: SEQ ID NOS: 170
SEQ ID NO 27
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                                                                                                                        3.0%; Score 13.4; DB 1; Length 17; 93.3%; Pred. No. 2.4e+02; 1ve 0; Mismatches 1; Indels
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; ORGANISM: Human immunodeficiency virus
US-10-157-580A-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             428 ACCCAGGACTCGGCT 442
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                                                                                                              3.0%
Best Local Similarity 93.3%
Matches 14; Conservative
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Best Local Similarity 80.0°
Matches 12; Conservative
                       ; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-232-634-12
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LENGTH: 17
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## APPLICANT: Kenneth, Debets

APPLICANT: Elisabeth, Roberts

TITLE OF INVENTION: OLIGONUCLECTIDE MEDIATED INHIBITION OF HEPATITIS B VIRUS AND HEPA

TITLE OF INVENTION: OLIGONUCLECTIDE MEDIATED INHIBITION OF HEPATITIS B VIRUS AND HEPA

FILE REFERENCE: 400/04208 (MBHBG2-249-E)

CURRENT APPLICATION NUMBER: US/10669, 841

CURRENT FILING DATE: 2003-09-23

PRIOR PILING DATE: 2003-09-23

PRIOR PELLING DATE: 2001-10-6-08

PRIOR PILING DATE: 2001-10-30

PRIOR PILING DATE: 2001-10-30

PRIOR PELLING DATE: 2001-10-30

PRIOR PELLING DATE: 2001-10-3-6

PRIOR PELLING DATE: 2000-10-3-6

PRIOR PELLING DATE: 2000-00-10

PRIOR PELLING DAT
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                                                                                       1; Indels
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                              Score 13.4; DB 1;
Pred. No. 2.4e+02;
); Mismatches 1;
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COTHER INFORMATION: oligonucleotide substrate US-10-669-841-3510
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                                                                                                                                                                                                                                                                                                                US-10-669-841-3510
Sequence 3510, Application US/10669841
Sequence 3510, US20040127446A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Lawrence, Blatt
                           ch 3.0%; Scc
1. Similarity 93.3%; Pre
14; Conservative 0;
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                                                                                                                                             30 GGTGGCCATTTTTG 44
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James, McSwiggen
David, Morrissey
                                                                                                                                                                                                   17 GATGGCCATTTTTG 3
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Patrice, Lee
Kenneth, Draper
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                              Query Match
Best Local Similarity
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Best Local Similarity
Matches 12; Conserv
                                                                                          Matches
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Sequence 4559, Application US/10287949A

Publication No. US20040102389A1

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: Bacobedo, Jaime

APPLICANT: Stinchcomb, Dan

APPLICANT: Escobedo, Jaime

TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor

FILE REFERENCE: MBHB00-876-N (400/049)

CURRENT APPLICATION NUMBER: US/10/287,949A

CURRENT PILING DATE: 2003-04-11

NUMBER OF SEQ ID NOS: 20822

SOFTWARE: Patentin version 3.0

LENGTH: 17
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APPLICANT: Pavco, Pam
APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: StinchCondb, Dan
APPLICANT: StinchCondb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00-876-N (400/049)
CURRENT APPLICATION NUMBER: US/10/287, 949A
CURRENT FILING DAME: 2003-04-11
NUMBER OF SEQ ID NOS: 2003-2
SOFTWARE: PatentIn version 3.0
SEQ ID NO 7501
LENGTH: 17
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CURRENT APPLICATION NUMBER: US/10/138,674
CURRENT FILING DATE: 2002-05-03
NUMBER OF SEQ ID NOS: 20822
SOFTWARE: PatentIn version 3.0
SEQ ID NO 7501
LENGTH: 17
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| Sequence 7501, Application US/10287949A
| Publication No. US20040102389A1
| GENERAL INFORMATION:
| APPLICANT: Ribozyme_Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    164 AGCAAACAAAATG 178
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Matches 13; Conservative
                                                                                                                                                                                                   ORGANISM: Homo sapiens
US-10-138-674-7501
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ORGANISM: Homo Bapiens
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TITLE OF INVENTION: NUCLEIC ACID BASED MODULATION OF VASCULAR ENDOTHELIAL GROWTH FACTY
TITLE OF INVENTION: NUCLEIC ACID BASED MODULATION OF VASCULAR ENDOTHELIAL GROWTH FACTY
FILE REPREBACE: REGERENCE: REGERENCE: REGERESIS RELATED DISEASES AND (
CURRENT APPLICATION NUMBER: US/0/12,633
CURRENT FILING DATE: 1095-10-26
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-0-1-08
PRIOR FILING DATE: 1995-0-1-08
PRIOR FILING DATE: 1995-0-1-08
PRIOR FILING DATE: 1995-0-1-07
PRIOR FILING DATE: 2000-11-07
PRIOR FILING DATE: 2001-10-7
PRIOR FI
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Publication No. US20050064495A1

GENERAL INFORMATION:

APPLICANT: Guida, Marco

FILE REFERENCE: DNA-20US2

CURRENT APPLICATION: WUMBER: US/10/949,004

CURRENT FILING DATE: 2004-09-23

PRIOR PELING DATE: 2002-08-30

PRIOR PLILING DATE: 2000-06-02

NUMBER OF SEQ ID NOS: 29

SOFTWARE: Patentin Ver. 2.1 and text editor

SEQ ID NO 11

LENGTH: 17
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Pred. No. 2.4e+02;
0; Mismatches 1; Indels
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US-10-949-004-12
; Sequence 12, Application US/10949004
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Best Local Similarity 93.3
Matches 14; Conservative
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US-10-712-633-490
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ORGANISM: Homo sapiens
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Matches 14; Conserv
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US-10-949-004-11
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                                       Sequence 6331, Application US/10669841
Sequence 6331, Application US/10669841
SEQUENCE 6331, Application US/10669841
SEQUENCE 6331, Application No. US20040127446A1
SEREAL INFORMATION:
APPLICANT: Lawrence. Blatt
APPLICANT: Dennis, Macajak
APPLICANT: Pamela, Pavco
APPLICANT: ON UNMER: US/10/669,941
CURRENT APPLICATION NUMBER: US 60/296,876
PRIOR FILING DATE: 2001-00-24
PRIOR PILING DATE: 2001-00-24
PRIOR APPLICATION NUMBER: US 60/333,055
PRIOR PILING DATE: 2001-01-24
PRIOR PILING PAPE: 2001-01-24
PRIOR PILING DATE: 2001-01-24
PRIOR PILING DATE: 2001-01-24
PRIOR PILING PAPE: 2001-01-24
NUMBER: OF SQ1/41,331
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OTHER INFORMATION: Description of Artificial Sequence: Nucleic Acid
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Publication No. US20040220128A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pamela
APPLICANT: Sandberg, Jennifer
APPLICANT: Gandberg, Jennifer
APPLICANT: Gozdon, Gilad
APPLICANT: McSwiggen, James
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ORGANISM: Artificial Sequence
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                               US-10-669-841-6231/c
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LENGTH: 17
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level)
TITLE OF INVENTION: RAS, HERZ and HIV
FILE REPERENCE: 400/046-102 (MHBB02-326-A)
CURRENT APPLICATION NUMBER: US/10/724,270
CURRENT PILING DATE: 2003-11-63
PRIOR FILING DATE: 2003-65-29
PRIOR FILING DATE: 2001-05-29
PRIOR FILING DATE: 2001-06-06
PRIOR PELICATION NUMBER: US 60/294,140
PRIOR PELICATION NUMBER: US 60/294,140
PRIOR FILING DATE: 2001-05-29
PRIOR FILING DATE: 2001-05-29
PRIOR FILING DATE: 2002-05-010
PRIOR PELICATION NUMBER: US 10/153,560
PRIOR FILING DATE: 2002-06-06
PRIOR PELICATION NUMBER: US 10/63,552
PRIOR PELICATION NUMBER: US 10/63,552
PRIOR PELICATION NUMBER: US 10/63,553
PRIOR PELICATION NUMBER: US 10/157,580
PRIOR PELICATION NUMBER: US 10/633,059
PRIOR PELICATION NUMBER: US 10/417,012
PRIOR PELICATION NUMBER: US 10/63-23
PRIOR PELICATION NUMBER: US 10/63-30
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APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level
TITLE OF INVENTION: RAS, HERZ and HIV
FILE REFERENCE: 400/046-US (MBHB02-326-A)
CURRENT APPLICATION NUMBER: US/10/724,270
CURRENT FILING DATE: 2003-11-26
PRIOR APPLICATION NUMBER: PCT/US/02/16840
PRIOR FILING DATE: 2002-05-29
PRIOR FILING DATE: 2002-05-29
PRIOR FILING DATE: 2001-09-10
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   Indels
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   Mismatches
                                                                                                                                                                                                                                                                                                                            Sequence 6679, Application US/10724270 Publication No. US20050080031A1 GENERAL INFORMATION:
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                                                                            428 ACCCAGGACTCGGCT 442
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   Conservative
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US-10-724-270-6679
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US-10-724-270-6679
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   12;
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Publication No. US20050080031A1

GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level
TITLE OF INVENTION: NUMBER: US/10/724,70
CURRENT APPLICATION NUMBER: US/10/724,70

PRIOR PELING DATE: 2001-05-10
PRIOR APPLICATION NUMBER: US 60/294,140
PRIOR PLING DATE: 2001-06-06
PRIOR APPLICATION NUMBER: US 60/294,140
PRIOR PLING DATE: 2001-06-06
PRIOR PELING DATE: 2002-06-06
PRIOR PLING DATE: 2002-06-06
PRIOR PLING DATE: 2002-06-06
PRIOR PLING DATE: 2002-06-06
PRIOR APPLICATION NUMBER: US 10/63,552
PRIOR PLING DATE: 2002-06-06
PRIOR APPLICATION NUMBER: US 10/63,559
PRIOR PLING DATE: 2002-06-29
PRIOR APPLICATION NUMBER: US 10/157,580
PRIOR PLING DATE: 2002-06-29
PRIOR APPLICATION NUMBER: US 10/63,552
PRIOR APPLICATION NUMBER: US 10/63,552
PRIOR APPLICATION NUMBER: US 10/63,553
PRIOR PLING DATE: 2002-06-29
PRIOR PLING DATE: 2002-06-29
PRIOR PRIOR PLING DATE: 2003-06-38
PRIOR PRIOR PLING DATE: 2003-06-38
PRIOR PLING DATE: 2003-06-38
PRIOR PRIOR PLING DATE: 2003-06-38
PRIOR PLING DATE: 2003-06-38
PRIOR PRIOR PLING DATE: 2003-06-38
                                                         APPLICANT: Guida, Marco
APPLICANT: Guida, Marco
APPLICANT: Hall, Jeff
JITLE OF INVENTION: GENETIC TYPING OF THE HUMAN CYTOCHROME P450 2A6 GENE
TITLE OF INVENTION: AND RELATED MATERIALS AND METHODS
FILE REPERENCE: DNA-20US2
CURRENT APPLICATION NUMBER: 10/212,634
PRIOR PILING DATE: 2004-09-23
PRIOR APPLICATION NUMBER: 10/212,634
PRIOR APPLICATION NUMBER: 10/212,634
PRIOR PILING DATE: 2000-06-02
PRIOR PILING DATE: 2000-06-02
NUMBER OF SEQ ID NOS: 29
SOFTWARE: Patentin Ver. 2.1 and text editor
SEQ ID NO 12
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NUMBER OF SEQ ID NOS: 6810
SOFTWARE: Patentin version 3.0
SEQ ID NO 6668
LENGTH: 17
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Pred. No. 2.4e+02;
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Publication No. US20050064495A1 GENERAL INFORMATION:
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Best Local Similarity 93.3%;
Matches 14; Conservative
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ORGANISM: Homo sapiens
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; ORGANISM: Homo sapiens
US-10-724-270-6668
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Best Local Similarity
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GENERAL INFORMATION:
APPLICANT: BCKer, David J.
APPLICANT: BCKer, David J.
APPLICANT: BCKer, David J.
APPLICANT: Bnomet, C. Frank
APPLICANT: Bnomet, C. Frank
APPLICANT: Hancak, Ronnie
APPLICANT: Hancak, Ronnie
APPLICANT: Hancak, Rounie
APPLICANT: APPLICANT: Anderson, Kevin
ITLE OF INVENTION: Modulation Of Telomere Length By Oligonucleotides Having A G-Core
ITLE OF INVENTION: Modulation Of Telomere Length By Oligonucleotides Having A G-Core
ITLE OF INVENTION: MOMBER: US/10/038,335
CURRENT FILING DATE: 2010-10-2
PRIOR APPLICATION NUMBER: 09/299,058
PRIOR APPLICATION NUMBER: 09/299,058
PRIOR PILING DATE: 1995-06-12
PRIOR PILING DATE: 1995-06-12
PRIOR PILING DATE: 1995-06-12
PRIOR PILING DATE: 1995-06-12
PRIOR PILING DATE: 1992-09-29
PRIOR PILING DATE: 1993-09-29
PRIOR PILING DATE: 1992-09-29
PRIOR PILING DATE: 1993-09-29
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Pred. No. 1.5e+02;
0; Mismatches 0; Indels
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2.9%; Score 13; DB 1; Lu
Best Local Similarity 100.0%; Pred. No. 1.5e+02;
Matches 13; Conservative 0; Mismatches 0;
                                 APPLICANT: Ahluwalia, Gurpreet S.
TITLE OF INVENTION: REDUCTION OF HAIR GROWTH
FILE REFERENCE: 00216-552001
CURRENT APPLICATION NUMBER: US/09/893,252
CURRENT FILING DATE: 2001-10-12
NUMBER OF SEQ ID NOS: 4
SOFTWARE: FastSEQ for Windows Version 4.0
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Best Local Similarity 100.0%; Pred. No. 1.5
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                                                                                                                                                                                                                                                                                                                     TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-10-038-335-1/c
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LENGTH: 13
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Publication No. US20050130186A1
GENERAL INFORMATION.
APPLICAMY: We Gene Technologies, Inc.
TITLE OF INVENTION: METHOD OF DETECTION CHIP AND FABRICATION METHOD TITLE OF INVENTION: DETECTION FILLS REPERSENCE: 12333-UG-PAA
CURRENT APPLICATION NUMBER: UP 92135134
PRIOR FILLING DATE: 2003-12-12
NUMBER OF SEQ ID NOS: 134
SOFTWARE: Patentin version 3.3
ERQ ID NO 41
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PRIOR APPLICATION NUMBER: US 60/296,249
PRIOR FILING DATE: 2001-06-06
PRIOR PELING DATE: 2001-05-29
PRIOR PAPLICATION NUMBER: US 60/294,140
PRIOR PELING DATE: 2001-05-29
PRIOR PELING DATE: 2002-09-10
PRIOR APPLICATION NUMBER: US 10/163,552
PRIOR PELING DATE: 2002-06-06
PRIOR PILING DATE: 2002-06-06
PRIOR PILING DATE: 2002-05-29
PRIOR PILING DATE: 2002-05-29
PRIOR PELING DATE: 2002-05-29
PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR PILING DATE: 2003-05-23
PRIOR PILING DATE: 2003-05-23
PRIOR PILING DATE: 2003-04-16
Remaining Prior Application data removed - See File Wrapper or PALM.
SOFTWARE: PatentIn version 3.0
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US-09-893-252-4/c
is Sequence 4, Application US/09893252
ipublication No. US20030012755A1
igeneral Information;
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Matches 12; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: Probe US-10-911-318-41
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US-10-724-270-6709
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Sequence 8, Application US/10463076;
Sequence 8, Application US/10463076;
Publication No. US2003021203241
GENERAL INFORMATION:
APPLICANT: Geron Corporation;
APPLICANT: Gryazov, Sergei
APPLICANT: Gryazov, Sergei
APPLICANT: Marray, Tracey
TITLE OF INVENTION: Oligonucleotide N3'-->P5' Thiophosphoramidates: Their Synthesis RILE REFERENCE: 039/004C
CURRENT APPLICATION NUMBER: US/10/463,076
CURRENT APPLICATION NUMBER: US 09/657,445
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 1999-09-10
PRIOR FILING DATE: 1999-09-10
PRIOR FILING DATE: 1999-10-19
NUMBER OF SEQ ID NOS: 9
SOFTWARE: Patentin version 3.1
SEQ ID NO 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; OTHER INFORMATION: Synthetic oligonucleotide with potential inhibition activity US-10-463-076-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           THER INFORMATION: Synthetic oligonucleotide with potential inhibition activity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence
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Best Local Similarity 100.1
Matches 13; Conservative
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13 TIGICTAACCCIA 1
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         APPLICANT: Wyatt, Jacquelline
APPLICANT: Hanecak, Ronnie
APPLICANT: Chiang, Ming-yi
APPLICANT: Sequence
TITLE OF INVENTION: Medulation Of Telomere Length By Oligonuclectides Having A G-Core
TITLE OF INVENTION: Sequence
FILE REFERENCE: ISIS-4976
CURRENT PRILING DATE: 2001-01-02
PRIOR APPLICATION NUMBER: 09/299,058
PRIOR APPLICATION NUMBER: 08/403,888
PRIOR PILING DATE: 1995-06-12
PRIOR PPLING DATE: 1993-09-29
PRIOR PILING DATE: 1993-09-29
PRIOR PILING DATE: 1993-09-29
PRIOR PILING DATE: 1992-09-29
PRIOR PILING DATE: 1992-09-29
PRIOR PILING DATE: 1992-09-29
NUMBER OF SEC ID NOS: 10
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Sequence 3, Application US/1025535

Publication No. US20030138814A1

GENERAL INFORMATION:
APPLICANT: Geron Corporation
APPLICANT: Granov, Sergai
APPLICANT: Tolman, Richard L.
APPLICANT: Morin, Gregg B.
TITLE OF INVENITON: Oligonuclectide Conjugates
FILE REFERENCE: 072/002P
CURRENT APPLICATION WUMBER: US/10/255,535
CURRENT FILING DATE: 2002-09-25
PRIOR FILING DATE: 2002-09-25
PRIOR FILING DATE: 2001-03-21
PRIOR FILING DATE: 2001-03-23
NUMBER OF SEQ ID NOS: 19
SEQ ID NO 3

LENGTH: 13
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    Wyatt, Jacqueline
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  46 CTAACCCTAACTG 58
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LENGTH: 13
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Page 120

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APPLICANT: Geron Corporation
APPLICANT: Graznov, Sergei
APPLICANT: Gryaznov, Sergei
APPLICANT: Gryaznov, Sergei
APPLICANT: Gryaznov, Krisztina
APPLICANT: Matray, Tracey
TITLE OF INVENTION OLIGonucleotide N3'-->P5' Thiophosphoramidates: Their Synthesis au
TITLE OF INVENTION OLIGONACCE 039,005C
CURRENT APPLICATION NUMBER: US,10/967,755
CURRENT APPLICATION NUMBER: US 10/463,076
PRIOR APPLICATION NUMBER: US 09/657,445
PRIOR APPLICATION NUMBER: US 60/153,201
PRIOR FILING DATE: 1999-09-10
PRIOR APPLICATION NUMBER: US 60/160,444
PRIOR FILING DATE: 1999-09-10
PRIOR APPLICATION NUMBER: US 60/160,444
PRIOR FILING DATE: 1999-10-19
NUMBER OF SEQ ID NOS: 9
SOFTWARE PATENTIN VEXEION 3.1
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GENERAL INFORMATION NO. US20050049408A1

APPLICANT: Geron Corporation

APPLICANT: Graznov, Sergei

APPLICANT: Matray, Tracey

TTLEOF INVENTION: Oligonuclectide N3'-->P5' Thiophosphoramidates: Their Synthesis and
FILE REPRENCE: 039/005C

CURRENT APPLICATION NUMBER: US/10/967, 755

CURRENT APPLICATION NUMBER: US 10/463, 076

PRIOR FILING DATE: 2003-66-17

PRIOR PELING DATE: 2003-60-09-08

PRIOR FILING DATE: 1999-09-10

PRIOR PLILING DATE: 1999-09-10

PRIOR FILING DATE: 1999-09-10

PRIOR FILING DATE: 1999-10-19

NUMBER OF SEQ ID NOS: 9

SOFTWARE: Patentin version 3.1

LENGTH: 13
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1.5e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.9%; Score 13; DB 100.0%; Pred. No. 1.5 tive 0; Mismatches
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Publication No. US20050049408A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
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Best Local Similarity 100.
Matches 13; Conservative
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Matches 13; Conservative
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                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 2
LENGTH: 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
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                                                                                                                                                                                                                  Sequence 18, Application US/10181823
Fublication No. US20040126752A1
Fublication No. US20040126752A1
GENERAL INFORMATION:
APPLICANT: Geron Corporation
APPLICANT: Gryaznov, Sergei
APPLICANT: Gryaznov, Sergei
APPLICANT: Schultz, Ronald
TITLE OF INVENTION: 2'-Arabino-Fluoroligonucleotide N3'-->P5' Phosphoramidates: Thei
TITLE OF INVENTION: Synthesis and Use
TITLE REFERENCE: 049/002
FILE REFERENCE: 049/002
CURRENT APPLICATION NUMBER: US/10/181,823
CURRENT FILING DATE: 2003-12-29
FRIOR FILING DATE: 2001-0119
FRIOR FILING DATE: 2001-0119
NUMBER: OF SEQ ID NOS: 23
SSOTID NO 18
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| Publication No. US20040126752A1
| Publication No. US20040126752A1
| Publication No. US20040126752A1
| Publication No. US20040126752A1
| APPLICANT: Geron Corporation
| APPLICANT: Geron Corporation
| APPLICANT: Granco, Sergei
| APPLICANT: Schultz, Ronald G
| TITLE OF INVENTION: 2'-Arabino-Fluoroligonucleotide N3'-->P5' Phosphoramidates: Thei
| TITLE OF INVENTION: Synthesis and Use
| TITLE OF INVENTION: Sy
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. 1.5e+02;
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2.9%; Score 13; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 1.5e+02;
Matches 13; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
2.9%; Score 13; DB
Best Local Similarity 100.0%; Pred. No. 1.5
Matches 13; Conservative 0; Mismatches
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; Sequence 2, Application US/10967755
   46 CTAACCCTAACTG 58
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13 TTGTCTAACCCTA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Homo sapiens
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                                                                                                                                                                                              US-10-181-823-18/c
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of Diseases or Conditions Related to Level
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Sequence 2188, Application US/20050080031A1

GENERAL INCORPATION:

MAPLICANT: McSwiggen, James

APPLICANT: McSwiggen, James

TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level

TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level

TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level

TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level

TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level

TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level

FILE REFREENCE: 400/046-US (MRH802-326-A)

CURRENT APPLICATION NUMBER: US 60/296, 249

PRIOR FILING DATE: 2001-06-06

PRIOR APPLICATION NUMBER: US 60/294, 140

PRIOR APPLICATION NUMBER: US 10/238,700

PRIOR FILING DATE: 2002-09-10

PRIOR PRIOR PLILING DATE: 2003-04-16

PRIOR APPLICATION NUMBER: US 10/44,853

PRIOR PRILING DATE: 2003-04-16

PRIOR APPLICATION NUMBER: US 10/417,012

PRIOR APPLICATION NUMBER: US 10/416

PRIOR APPLICATION NUMBER: US 10/417,012

PRIOR APPLICATION NUMBER: US 10/416

PRIOR PRIOR PRIOR DATE: 2003-04-16

PRIOR PRIO
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                           APPLICANT: McSwiggen, James
TITLE OF INVENTION: Nucleic Acid Treatment of
FILE REPERENCE: 400/057 (MBHB01-1158-A)
CURRENT APPLICATION NUMBER: US/10/238,700
CURRENT FILING DATE: 2002-09-18
PRIOR APPLICATION NUMBER: PCT/US 02/16840
PRIOR FILING DATE: 2002-05-29
PRIOR PILING DATE: 2002-05-29
PRIOR FILING DATE: 2001-09-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.9%; Score 13; DB 100.0%; Pred. No. 2.6 iive 0; Mismatches
      APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 4666
SOFTWARE: PatentIn version 3.0
SEQ ID NO 3509
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            225 CCTGCCCAGCCCC 237
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            225 CCTGCCCAGCCCC 237
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Best Local Similarity 100.0
Matches 13; Conservative
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Best Local Similarity 100.
Matches 13, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13 cereceasecee 1
                                                                                                                                                                                                                                                                                                                                                                                                                  ; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-238-700-3509
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US-10-724-270-2188
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                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: A POLYMUCLEIC ACID ISOLATED FROM A
PORCINE REPRODUCTIVE AND RESPIRATORY SYNDROME VIRUS (PRRSV)
A PROTEIN ENCODED BY THE POLYMUCLEIC ACID, A VACCINE
PREPARED FROM OR CONTAINING THE POLYMUCLEIC ACID OR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ò
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 77
CORRESPONDENCE ADDRESS:
ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
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STRBET: 1755 S. Jefferson Davis Highway, Suite 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
COMPUTER: DEADELS
CONTUTER: DEADELS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/10/618,779
FILING DATE: 15-Jul-2003
CIASSIFICATION ADATA:
PROPLICATION NUMBER: US/08/301,435
FILING DATE: 01-SEP-1994
APPLICATION NUMBER: US 08/131,625
FILING DATE: 05-OCT-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 2.9%; Score 13; DB 1; Length 16; Best Local Similarity 100.0%; Pred. No. 2.3e+02; Matches 13; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Lavalleye, Jean-Paul M.P.
REGIGYRATION NUMBER: 31,451
REFERENCE/DOCKET NUMBER: 4625-021-55X CIP
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
SEQUENCE DESCRIPTION: SEQ ID NO: 34:
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8.10-238-700-3509/c
; Sequence 3509, Application US/10238700
; Publication No. US20030153521A1
                                                                                                                                                                                                                                                                                                MENG, XIANG-JIN
HALBUR, PATRICK G.
MOROZOV, IGOR
LUM, MELISSA A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (703) 413-3000
                                                                                                                                                                            Sequence 34, Application US/10618779
Publication No. US20050175633A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (703) 413-2220
TELEX: 248855 OPAT UR
INFORMATION FOR SEQ ID NO: 34:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: unknown
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                                                                                                                                                                                                                                                                  APPLICANT: PAUL, PREM S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       268 GGGCTTCTCCGG 280
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Arlington
STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: U.S.A.
13 CTAACCCTAACTG
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                                                                                                                    RESULT 308
US-10-618-779-34
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APPLICANT: Stindhoomb, Dann
TITLE OF INVENTION: Method and Reagent for the Inhibition of Telomerase Enzyme
FILE REPERENCE: MEHBOO-882-(400/019)
CURRENT APPLICATION NUMBER: US/10/712,672
CURRENT FILING DATE: 2003-11-13
PRIOR APPLICATION NUMBER: US/09/653,225
PRIOR PILING DATE: 2000-09-14
PRIOR PILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/197,769
PRIOR APPLICATION NUMBER: 60/197,769
PRIOR APPLICATION NUMBER: 60/197,769
PRIOR APPLICATION NUMBER: 60/190,713
PRIOR PILING DATE: 1999-08-31
NUMBER OF SEQ ID NOS: 5586
SOFTWARE: PARENTLING DATE: 1999-08-31
NUMBER OF SEQ ID NOS: 5586
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 61, Application US/10483958

Publication No. US20040254363A1

GENERAL INFORMATION:

APPLICANT: PRICE FOUNDATION LIMITED

APPLICANT: PRICE FOUNDATION LIMITED

APPLICANT: PREASE, Mercedith

APPLICANT: STAGER, Mercedith

TITLE OF INVENTION: GENES AND SNPS ASSOCIATED WITH EATING DISORDERS

CURRENT FILING DATE: 2004-01-6

PRIOR PELING DATE: 2004-01-6

PRIOR PELING DATE: 2002-07-16

PRIOR PELING DATE: 2001-07-16

PRIOR APPLICATION NUMBER: US 60/305,153

PRIOR PELING DATE: 2001-07-16

PRIOR PELING DATE: 2001-07-16

PRIOR PELING DATE: 2001-07-16

PRIOR PELING DATE: 2001-11-13

PRIOR PELING DATE: 2001-11-13

PRIOR PELING DATE: 2001-11-13

PRIOR PELING DATE: 2001-11-19

PRIOR PELING DATE: 2001-11-19

PRIOR PELING DATE: 2001-11-19

PRIOR PELING DATE: 2001-12-19

PRIOR PELING DATE: PERCENTIN VERSION 3.1

PRIOR PELING DATE: PERCENTIN VERSION 3.1

PRIOR PERCENTIN VERSION 3.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; OTHER INFORMATION: HTR1D probe: FAM and MGB tagged US-10-483-958-61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     247 CCTGGAGGCCGCGGTC 262
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 ccuggaggccgcagcc 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 GGTTGCGGAGGGTGGG 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-712-672-1787
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Best Local Similarity
Watches 14; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 1787
LENGTH: 16
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81.2%; Pred. No. 2.4e+02;
iive 1; Mismatches 2; Indels
                                                                                                                   Sequence 159, Application US/08887505
Fublication No. US20020081577A1
Fublication No. US20020081577A1
GENERALI INFORMATION:
APPLICANT: Kilkuskie, Robert E.
APPLICANT: Goodelid, John
APPLICANT: Wolfe, Jia L.
APPLICANT: Wolfe, Jia L.
APPLICANT: Roberts, Perer C.
APPLICANT: Roberts, No. US20020081577A11 A.
APPLICANT: Roberts, No. US20020081577A11 A.
APPLICANT: Whenty A.
APPLICANT: Wolfer, Jia L.
APPLICANT: W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 02109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATE: PATENTER: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
APPLICATION NUMBER: US/08/887,505
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:

CLASSIFICATION: 514

PRIOR APPLICATION: 514

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/471,968

FILING DATE: 06-JUN-1995

ATTONNEY/AGENT INPORMATION:

NAME: KATHER, AND-LOUISE

REGISTRATION NUMBER: 33,523

REFERENCE/DOCKET NUMBER: HYZ-040CIP

TELECOMMUNICATION INPORMATION:

TELECOMMUNICATION INPORMATION:

TELEFAX: (617) 526-6000

TELEFAX: (617) 526-5000

INFORMATION FOR SEQ ID NO: 159:

SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs

TYPE: nucleic acid

TYPE: nucleic acid
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APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Chowrira, Bharat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   199 CCCTCCCGGGGACCTG 214
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13 CCTGCCCAGCCCC 1
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Best Local Similarity 81.2
Matches 13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
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STATE: MA
COUNTRY: USA
                                                                                                                                                                                             US-08-887-505-159
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US-10-712-672-1787
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US-08-887-505-159
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RESULT 314

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Pred. No. 2.7e+02;
0; Mismatches 2;
                                                         PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 08/679,645
FILING DATE: July 12, 1996
APPLICATION NUMBER: 60/001,135
FILING DATE: July 13, 1995
APPLICATION NUMBER: 08/300,726
FILING DATE: September 2, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 219/247
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/961,077
FILING DATE: 21-Sep-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE DESCRIPTION: SEQ ID NO: 828:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-780-533A-58/c
; Sequence 58, Application US/09780533A
                                                                                                                                                                                                                                                                    TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                     TELEX: 67-3510 INFORMATION FOR SEQ ID NO: 828:
                                                                                                                                                                                                                                                                                                                                    LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            107 GCTGACTTTCAGCGGG 122
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SOFTWARE: Patentin version 3.0
SEQ ID NO 27
LENGTH: 17
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                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
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Best Local Similarity 62.5
Matches 10; Conservative
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Best Local Similarity 87.5
Matches 14; Conservative
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US-09-780-533A-27
                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-961-077-828
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2.8%; Score 12.8; DB 1; Length 16;
Best Local Similarity 87.5%; Pred. No. 2.4e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Folkerts, Otto
Merlo, Donald J.
TITLE OF INVENTION: COMPOSITION AND METHODS FOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 828, Application US/09961077
Publication No. US20030014775A1
GENERAL INFORMATION: Michael G.
APPLICANT: Zwick, Michael G.
Edington, Brent E.
McSwiggen, James A.
McNiggen, Lining
Skokut, Thomas A.
Young, Scott A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DOS 5.0
         NUMBER OF SEQUENCES: 1263
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                             ) OTHER INFORMATION: Template sequence US-10-730-771-444
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
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US-09-877-478-1053
                                                                                                                                                                     TYPE: RNA
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                     APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: Chowrira, Bharat
APPLICANT: Chowrira, Bharat
APPLICANT: Haeberli, Pete
ITITE OF INVENTION: Method and Reagent for the Inhibition of NOGO Gene
FILE REFERENCE: MBHB00, 878-A, (400/011)
CURRENT APPLICATION NUMBER: US/09/780,533A
CURRENT FILING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 6679
SOCTWARE: Patentin version 3.0
SEQ ID NO 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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nilarity 87.5%; Pred. No. 2.7e+02;
Conservative 0; Mismatches 2; Indels
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Sequence 1810, Application US/09780533A
Publication No. US20030060611A1
SERERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, Jim
APPLICANT: Chowaira, Bharat
APPLICANT: Chowaira, Bharat
APPLICANT: Haeberli, Pete
                                                                                                                                                                                                                                                                                                                                                                                                    363 GCCGCAGGAAGAGAA 378
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Publication No. US20030060611A1
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                                                                                                                                                                                                                                                                                          ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: RNA
CORGANISM: Homo sapiens
US-09-780-533A-59
                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                                                                                                       US-09-780-533A-58
                                                                                                                                                                                                                                                                      TYPE: RNA
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RESULT 320

Sequence 1053, Application US/09877478

Sequence 1053, Application US/09877478

Sequence 1053, Application US/09877478

Publication No. US20030068301A1

APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Reckiggen, Jim
APPLICANT: Blatt, Larry

APPLICANT: Mexidgen, Jim
APPLICANT: Morrisaey, Dave
ITITLE OF INVENTION: Method and Reagent for Inhibiting Hepatitis B Virus Replication
FILE REFERENCE: MBHBO-045-H (400/029)
CURRENT APPLICATION NUMBER: US 09/636, 385
FRIOR FILING DATE: 2000-03-20

FRIOR APPLICATION NUMBER: US 09/531,025
FRIOR FILING DATE: 2000-03-20

FRIOR APPLICATION NUMBER: US 09/534,025
FRIOR FILING DATE: 1999-02-10
FRIOR PLING DATE: 1999-02-04
FRIOR FILING DATE: 1999-02-04
FRIOR FILING DATE: 1999-02-05
FRIOR FILING DATE: 1999-02-06
FRIOR FILING DATE: 1999-02-06
FRIOR APPLICATION NUMBER: US 08/434,504
FRIOR FILING DATE: 1999-11-08
FRIOR FRIOR PROPERTY 11-07
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TITLE OF INVENTION: Method and Reagent for the Inhibition of NOGO Gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 2.8%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 2.7e+02; Matches 14; Conservative 0; Mismatches 2; Indels
                     FILE REFERENCE: MBHBOO, 878-A (400/011)
CURRENT APPLICATION NUMBER: US/09/780,533A
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: US 60/181,797
PRIOR FILING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 6679
SEQ ID NO 1810
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; Sequence 1036, Application US/09848754A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; ORGANISM: Homo sapiens
US-09-780-533A-1810
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US09436060A-16.rnpb.sl

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US-05-848-754A-1039/c
| Sequence 1039, Application US/09848754A
| Sequence 1039, Application US/09848754A
| Sequence 1039, Application O. US20030073207A1
| Sequence 1039, Application No. US20030073207A1
| FUBLICANT: Ribozyme Pharmaceuticals, Inc.
| APPLICANT: Ribozyme Pharmaceuticals, Inc.
| TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
| TITLE OF INVENTION: Levels Of Epidermal Growth Factor Receptors
| FILE REFERENCE: MHBB00-958-I (400/018)
| CURRENT APPLICATION NUMBER: US/09/848,754A
| CURRENT FILING DATE: 2001-05-03
| SOFTWARE: PatentIn version 3.0
| SEQ ID NO 1039
| TENGRAL 1039
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Publication No. US20030073207A1

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc.

TITLE OF INVENTION: Levels of Epidermal Growth Factor Receptors

TITLE REFERENCE: MBHB00-958-I (400/018)

CURRENT APPLICATION NUMBER: US/09/848,754A

CURRENT FILING DATE: 2001-05-03

NUMBER OF SEQ ID NOS: 9645

SEQ ID NO 1653
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87.5%; Pred. No. 2.7e+02;
iive 0; Mismatches 2; Indels
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87.5%; Pred. No. 2.7e+02;
tive 0; Mismatches 2;
87.5%; Pred. No. 2.76 ive 0; Mismatches
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Publication No. US20030087847A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                          199 CCCTCCCGGGGACCTG 214
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Holman, Patricia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; ORGANISM: Homo sapiens
US-09-848-754A-1039
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US-09-848-754A-1653
Best Local Similarity
Matches 14; Conserv
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Matches 14; Conser
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US-09-776-474-7/c
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APPLICANT:
APPLICANT:
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                                         APPLICANT: Ribozyme Pharmaceuticals, Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Levels of Epidermal Growth Factor Receptors
CURRENT SPERIENCE: MBHB00-958-I (400/018)
CURRENT APPLICATION NUMBER: US/09/848,754A
SURRENT FILLING DATE: 2001-05-03
NUMBER OF SEQ ID NOS: 9645
SOFTWARE: Patentin version 3.0
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Publication No. US20030073207A1
Publication No. US20030073207A1
ARDELIANZI INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Levels of Epidermal Growth Factor Receptors
FILE REFERENCE: MBHB00-958-I (400/018)
CURRENT APPLICATION NUMBER: US/09/848,754A
NUMBER OF SEQ ID NOS: 9645
SOFTWARE: Patentin version 3.0
SEQ ID NO 1037
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Sequence 1038, Application US/09848754A

Publication No. US20030073207A1

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc.

TITLE OF INVENTION: Levels of Epidermal Growth Factor Receptors

FILE REFERENCE: MBHB00-958-I (400/018)

CURRENT APPLICATION NUMBER: US/09/848,754A

CURRENT PILING DATE: 2001-05-03

NUMBER OF SEQ ID NOS: 9645

SOFTWARE: Patentin version 3.0
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Publication No. US20030073207A1
GENERAL INFORMATION:
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CORGANISM: Homo sapiens
US-09-848-754A-1037
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US-09-848-754A-1036
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US-09-848-754A-1037/c
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LENGTH: 17
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LENGTH: 17
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Sequence 335, Application US/09930423
Publication No. US20030092003A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: Mibozyme Pharmaceuticals, Inc.
TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease;
FILE REPERENCE: MBHB00, 918-A 400/027
CURRENT APPLICATION NUMBER: US/09/930,423
CURRENT PILING DATE: 2001-08-15
NUMBER OF SEQ ID NOS: 4553
SOFTWARE: PatentIn version 3.0
SOFTWARE: PatentIn version 3.0
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CURRENT APPLICATION NUMBER: US/09/930423

FILE REFERENCE: MEMBOO, 198, 400/027

CURRENT APPLICATION NUMBER: US/09/930,423

CURRENT APPLICATION NUMBER: US/09/930,423

NUMBER OF SEQ ID NOS: 4553

SOFTWARE: Patentin version 3.0

SEQ ID NO 1159
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2.8%; Score 12.8; DB 1;
Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2;
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                                                                                                        Query Match
Best Local Similarity 87.5'
Matches 14; Conservative
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Matches 14; Conservative
                                      ; ORGANISM: Homo Sapiens
US-09-930-423-334
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TYPE: RNA
CRGANISM: Homo Sapiens
US-09-930-423-335
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CORGANISM: Homo Sapiens
US-09-930-423-1159
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US-09-930-423-1470
                     TYPE: RNA
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APPLICANT: Fattaey, Ali
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: McEvagen, Jim
TITLE OF INVENTION: McEvagen, Jim
TITLE OF INVENTION: McEvagent for the Inhibition of Checkpoint Kinase-1 (CHK
TITLE OF INVENTION: McEvagen
TITLE OF INVENTION: McEvagen
TITLE OF INVENTION: McEvagen
TITLE OF INVENTION: McMark: Mc10/008)
CURRENT APPLICATION NUMBER: US/09/776,474
CURRENT FILING DATE: 2001-00-02
PRIOR FILING DATE: 2000-03-02
RICH APPLICATION NUMBER: US 60/179,983
RICH FILING DATE: 2000-03-02
NUMBER OF SEQ ID NOS: 2992
SOFTWARE: PatentIn version 3.0
ERNGTH: 17
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Publication No. US20030092003A1
Publication No. US20030092003A1
Publication No. US20030092003A1
APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: Machod and Reagent for the Treatment of Alzheimer's Disease
FILE REPERENCE: MBHB00,918-A 400/027
CURRENT PRILING DATE: 2001-08-15
NUMBER OF SEQ ID NOS: 4553
ASSPTWARE: Patentin version 3.0
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Publication No. US20030092003A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: Match and Reagent for the Treatment of Alzheimer's Disease
TITLE OF INVENTION: Mchod and Reagent for the Treatment of Alzheimer's Disease
CURRENT FILING DATE: 2001-08-15
NUMBER OF SEQ ID NOS: 4553
SOFTWARER PATCH NOT SEQ ID NOS: 4553
SOFTWARER PATCH NOT SEQ ID NOS: 4553
LENGTH: 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     272 CITCTCCGGAGGCACC 287
                                                                                                                                                                                                                                                                                                   TYPE: RNA
ORGANISM: Artificial Sequence
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CORGANISM: Homo Sapiens
US-09-930-423-333
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US-09-930-423-334/c
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LENGTH: 17
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Sequence 3639, Application US/09740332
| Publication No. US20030125270A1
| Publication No. US20030125270A1
| GENERAL INFORMATION:
| TITLE OF INVENTION: Braymatic Nucleic Acid Treatment of Diseases or Conditions Relate:
| TITLE OF INVENTION: Hepatitis C Virus Infection |
| TITLE OF INVENTION: Hepatitis C Virus Infection |
| TITLE OF INVENTION: Hepatitis C Virus Infection |
| TITLE OF INVENTION: Hepatitis C Virus Infection |
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| TITLE OF INVENTION: Hepatitis C Virus Infection |
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| TITLE OF INVENTION: Hepatitis C Virus Infection |
| TITLE OF INVENTION: Hepatitis C Virus Infection |
| TITLE OF INVENTION: Hepatitis C VIII |
| TITLE OF INVENTION: Hepatitis C VIII |
| TITLE OF INVENTION INVENTION |
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Publication No. US20030125270A1

GENERAL INFORMATION:

APPLICATY: RIDOZYME Pharmaceuticals Inc.

TITLE OF INVENTION: Hepatitis C Virus Infection

STATUS OF THE REFERENCE INVENTION: Hepatitis C Virus Infection

STATUS OF THE STATUS INVENTION: Hepatitis C Virus Infection

STATUS OF THE STATUS INVENTION: Hepatitis C Virus Infection

STATUS OF THE STATUS INVENTION: Hepatitis C VIRUS INVENTION IN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.8%; Score 12.8; DB 1; 75.0%; Pred. No. 2.7e+02; tive 2; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 2.8%; Score 12.8; DB 1; Best Local Similarity 87.5%; Pred. No. 2.7e+02; Matches 14; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                    ; OTHER INFORMATION: oligonucleotide substrate US-09-740-332-1425
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: oligonucleotide substrate US-09-740-332-3639
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LOCATION: OTHER INFORMATION: oligonucleotide substrate
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    SOFTWARE: Patentin version 3.0
SEQ ID NO 1425
LENGTH: 17
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ORGANISM: artificial sequence
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                                                                                                                                        TYPE: RNA
ORGANISM: artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16 ecacerecededere 1
                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 75...
Best Local Similarity 75...
                                                                                                                                                                                                                                                                   NAME/KEY: misc_feature LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: misc_feature
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                                                                                                                                                                                                                              FEATURE:
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US-09-740-332-1425

US-09-740-332-1425

Sequence 1425, Application US/09740332

Sequence 1425, Application US/09740332

Publication No. US20030125270A1

GENERAL INFORMATION:

APPLICANT: RIDEXTME Pharmaceuticals Inc.

TITLE OF INVENTION: Hepatitis C Virus Infection

FILE REFERENCE: RPI 400/003

FILE REFERENCE: RPI 400/003

CURRENT APPLICATION UMBER: US/09/740,332

CURRENT FILING DATE: 2001-03-26

NUMBER OF SEQ ID NOS: 9704
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TITLE OF INVENTION: Method and Reagent for the Inhibition of NOGO and NOGO Receptor (
FILE REFERENCE: MBHB00-878-C (400/017)
CURRENT FILING NOW NOWBER: US/09/827,395A
CURRENT FILING DATE: 2001-04-05
PRIOR APPLICATION NUMBER: 09/780,533
PRIOR FILING DATE: 2000-02-19
PRIOR FILING DATE: 2000-02-11
NUMBER: OF SEQ ID NOS: 2617
SOFTWARE: Patentin version 3.0
SEQ ID NO 899
                                                                                                                                                                   APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Mcthod and Reagent for the Treatment of Alzheimer's Disease
FILE REFERENCE: MBHB00,918-A,400/027
CURRENT APPLICATION WHERE: US/09/930,423
CURRENT FILING DATE: 2001-08-15
NUMBER OF SEQ ID NOS: 4553
SEQ ID NO 1470
LENGTH: 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 17;
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Best Local Similarity 75.0%; Pred. No. 2.7e+02;
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Publication No. US20030113891A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaccuticals, Inc.
APPLICANT: Lawrence Blatt
                                                                                                                                        APPLICANT: Ribozyme Pharmaceuticals, Inc
Sequence 1470, Application US/09930423
Publication No. US20030092003A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  263 GGCCGGGGCTTCTCC 278
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; ORGANISM: Homo Sapiens
US-09-930-423-1470
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US-09-827-395A-899
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Sequence 334, Application US/09745237A
Publication No. US20030143708A1
GENERAL INFORMATION:
APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: Machod and Reagent for the Treatment of Alzheimer's Disease
TITLE OF INVENTION: MCHOD and Reagent for the Treatment of Alzheimer's Disease
CURRENT APPLICATION NUMBER: US/09/745,237A
CURRENT FILING DATE: 2002-04-15
NUMBER OF SEQ ID NOS: 4550
SOFTWARE: PatentIn version 3.0
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US-09-745-237A-1159/C
Sequence 1159, Application US/09745237A
Sequence 1159, Application US/09745237A
Sequence 1159, Application US/09043708A1
Sequence 1159, Application No. US20030143708A1
GENERAL INFORMATION:
APPLICANT: Blact, Larry
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Treatment of Alzheimer's Disease;
FILE REPERENCE: 400/007 (MBHBO0-918-A)
CURRENT APPLICATION NUMBER: 2002-04-15
CURRENT FILING DATE: 2002-04-15
SOFTWARE: PatentIn version 3.0
SEQ ID NO 1159
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Publication No. US20030143708A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
TILLE OF INVENTION: McHod and Reagent for the Treatment of Alzheimer's Disease
FILE REPERENCE: 400/007 (MBHB00-9.18-A)
CURRENT APPLICATION NUMBER: US/09/745,237A
CURRENT PILING DATE: 2002-04-15
NUMBER OF SEQ ID NOS: 4550
SOFTWARE: PatentIn version 3.0
SEQ ID NO 335
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US-09-745-237A-334
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: RNA
ORGANISM: Homo sapiens
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                       ) Sequence 331, Application US/09792818
) Publication No. US20030134806A1
) GENERAL INFORMATION:
| APPLICANT: Ribozyme Pharmaceuticals, Inc.
| APPLICANT: Jarvis, Thale
| APPLICANT: Von Carlowitz, Ira
| APPLICANT: Won Carlowitz, Ira
| APPLICANT: Hamblin, Paul
| APP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 333, Application US/09745237A
Publication No. US20030143708A1
GENERAL INFORMATION.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: Method and Reagent for the Treatment of Alzheimer's Disease
FILE REPERENCE: 400/007 (MBHB00-918-A)
CURRENT APPLICATION NUMBER: US/09/745,237A
CURRENT APPLICATION NUMBER: US/09/745,237A
CURRENT APPLICATION NUMBER: US/09/745,237A
CURRENT APPLICATION NUMBER: US/09/745,237A
CURRENT PILING DATE: 2002-04-15
NUMBER OF SEQ ID NOS: 4550
SOFTWARE: Patentin version 3.0
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                                                            Ouery Match 2.8%; Score 12.8; DB 1; Length 17; Best Local Similarity 87.5%; Pred. No. 2.7e+02; Matches 14; Conservative 0; Mismatches 2; Indels
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; ORGANISM: Homo sapiens
US-09-745-237A-333
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US-09-745-237A-333/c
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US-09-740-332-3755
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Sequence 3639, Application US/09817879

Publication No. US20030171311A1

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals Inc.

TITLE OF INVENTION: Hepatitis C Virus Infection

TITLE OF INVENTION INVENTION
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Sequence 3755, Application US/09817879

Sequence 3755, Application US/09817879

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Sequence 3755, Application US/09817879

Fublicant: Nicoration Nicoration US/09/09/09/C

TITLE OF INVENTION: Hepatitie C Virus Infection

TITLE OF INVENTION: Hepatitie C Virus Infection

FILE REFERENCE: MBH800-801-F

CURRENT APPLICATION NUMBER: US/09/817,879

CURRENT PILING DATE: 2001-03-26

SOFTWARE: Patentin version 3.0

SEQ ID NO 3755
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87.5%; Pred. No. 2.7e+02;
tive 0; Mismatches 2;
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; Sequence 738, Application US/10060895A
; Publication No. US20030104403A1
; APPLICANT: Zhang, Jian
; APPLICANT: Gu, Yizhong
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ORGANISM: artificial sequence
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ORGANISM: artificial sequence
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Best Local Similarity 87.51
Matches 14; Conservative
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                                                                                                                    US-09-817-879-3639/c
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Publication No. US20030171311A1

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals Inc.

TITLE OF INVENTION: Hepatitis C virus Infection

TITLE OF INVENTION

TITLE
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Publication No. US20030143708A1
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: Macada and Reagent for the Treatment of Alzheimer's Disease FILE REFERENCE: 400/007 (MBHB00-918-A)
CURRENT APPLICATION NUMBER: US/09/745,237A
CURRENT FILING DATE: 2002-04-15
NUMBER OF SEQ ID NOS: 4550
SOFTWARE: Patentin version 3.0
SEQ ID NO 1470
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2.8%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                       12 GGTGGCCTGGGAGGG 27
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Matches 14; Conservative
                                  TYPE: RNA
GRGANISM: Homo sapiens
US-09-745-237A-1159
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ORGANISM: Homo sapiens
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US-09-745-237A-1470
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US-09-817-879-1425
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LENGTH: 17
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APPLICANT: MCSWiggen, Jim
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Nucleic acid treatment of diseases or conditions related to level:
TITLE OF INVENTION: HERS.
TITLE OF INVENTION: HERS.
CURRENT APPLICATION NUMBER: US/10/163,552
CURRENT FILING DATE: 2002-06-06
NUMBER OF SEQ ID NOS: 1997
SOFTWARE: Patentin version 3.0
SEQ ID NO 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 869, Application US/10163552
Publication No. US20030105051A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Roswiggen, Jim
TITLE OF INVENTION: Nucleic acid treatment of diseases or conditions related to level
TITLE OF INVENTION: HER2
FILE REFERENCE: MBHB01-1653-A (400/014)
CURRENT APPLICATION NUMBER: US/10/163,552
CURRENT FILING DATE: 2002-06-06
NUMBER OF SEQ ID NOS: 1997
SOFTWARE: PatentIn version 3.0
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2.8%; Score 12.8; DB 1;
Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2;
                       PRIOR FILING DATE: 2001-08-30
NUMBER OF SEQ ID NOS: 1682
SOFTWARE Acomica Sequence Listing Engine
SEQ ID NO 739
LENGTH: 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16 GCGCGGATCAGGCCTT 1
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                                                                                                                                                                                                       TYPE: DNA
CORGANISM: Homo sapiens
US-10-060-895A-739
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-163-552-20
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ORGANISM: Homo sapiens
US-10-163-552-869
                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 14; Conserv
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US-10-163-552-869/c
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LENGTH: 17
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US-10-060-895A-739/C

US-10-060-895A-739/C

US-10-060-895A-739/C

US-10-060-895A-739/C

US-10-060-895A-739/C

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87.5%; Pred. No. 2.7e+02;
iive 0; Mismatches 2; Indels
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                                                                                                                                             CURRENT FILING DATE: 2002-06-10

PRIOR APPLICATION NUMBER: PCT/USO1/00666

PRIOR APPLICATION NUMBER: PCT/USO1/00667

PRIOR PILING DATE: 2001-01-30

PRIOR PILING DATE: 2001-01-30

PRIOR PILING DATE: 2001-01-30

PRIOR PLICATION NUMBER: PCT/USO1/00669

PRIOR PILING DATE: 2001-01-30

PRIOR PILING DATE: 2001-01-30

PRIOR PILING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: PCT/USO1/00665

PRIOR PILING DATE: 2001-01-30

PRIOR PILING DATE: 2001-05-23

PRIOR PILING DATE: 2001-01-30

PRIOR PILING DATE: 2001-01-30
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Best Local Similarity 87.5
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
ORGANISM: Homo sapiens
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APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, James
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level
FILE REFERENCE: 400/057 (MBHB01-1158-A)
CURRENT APPLICATION NUMBER: US/10/238,700
CURRENT FILING DATE: 2002-09-18
PRIOR FILING DATE: 2002-05-29
PRIOR FILING DATE: 2001-09-10
NUMBER OF SEQ ID NOS: 4666
SOFTWARE: Patentin version 3.0
SEQ ID NO 2821
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Publication No. US20030153521A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: MCSWiggen, James
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level
FILE REPERENCE: 400/057 (MRHB01-1158-A)
CURRENT APPLICATION NUMBER: US/10/238,700
CURRENT FILING DATE: 2002-09-18
PRIOR FILING DATE: 2002-09-18
PRIOR FILING DATE: US 60/318,471
PRIOR FILING DATE: 2001-09-10
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                                                                                                                                                                                                                                                                                                              2; Indels
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87.5%; Pred. No. 2.7e+02;
trive 0; Mismatches 2;
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Pred. No. 2.7e+02;
2; Mismatches 2;
                             60/318,471
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. Sequence 2821, Application US/10238700
. Publication No. US20030153521A1
. GENERAL INFORMATION:
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PRIOR FILING DATE: 2002-05-29
PRIOR APPLICATION NUMBER: US 60/
PRIOR FILING DATE: 2001-09-10
NUMBER OF SEQ ID NOS: 4666
SOFTWARE: Patentin version 3.0
SEQ ID NO 2700
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SOFTWARE: PatentIn version 3.0
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Best Local Similarity 75.0%;
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Best Local Similarity 87.55
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12; Conservative
                                                                                                                                                                    ; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-238-700-2700
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; ORGANISM: Homo sapiens
US-10-238-700-2821
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ORGANISM: Homo sapiens
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LENGTH: 17
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                                                               RESULT 349

US-10-156-306-4930

US-10-156-306-4930

Sequence 4990, Application US/10156306

Publication No. US2030319017A1

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: MCSWiggen, James

TITLE OF INVENTION: Levels of IKK-Gamma and PKR

TITLE OF INVENTION: Levels of IKK-Gamma and PKR

FILE REFERENCE: MBHB01-664-A (400/050)

CURRENT APPLICATION NUMBER: US/10/156,306

CURRENT FILING DATE: 2002-05-28

SOFTWARE: PatentIn version 3.0

SEQ. 10 V 930
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Publication No. US2003011901741
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, James
TITLE OF INVENTION: Levels of IKK-Gamma and PKR
FILE REFERENCE: MBHB01-664-A (400/050)
CURRENT APPLICATION NUMBER: US/10/156,306
CURRENT FILING DATE: 2002-05-28
NUMBER OF SEQ ID NOS: 8013
SOFTWARE: PatentIn version 3.0
FERSIOL DATE: SEQ ID NOS: 8013
SOFTWARE: PatentIn version 3.0
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Publication No. US20030153521A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, James
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level
FILE REFERENCE: 400/057 (MBHB01-1158-A)
CURRENT APPLICATION NUMBER: US/10/238,700
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                249 TGGAGGCCGCGGTCGG 264
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    17 GAGGAGGGTGGGCTG
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Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: RNA
CORGANISM: Homo sapiens
US-10-156-306-4930
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: RNA
ORGANISM: Homo sapiens
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US-10-238-700-2700/c
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US-10-156-306-6942
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Sequence 1249, Application US/10230006

Bublication No. US20030191077A1

GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Rosnaugh, Kathy
APPLICANT: Fornaugh, Kathy
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: METHOD AND REAGENT FOR THE TREATMENT OF ASTHWA AND ALLERGIC CONDITILE REFERENCE: 400/056 (MBHB01-1110)
CURRENT APPLICATION NUMBER: US/10/230,006
CURRENT FILING DATE: 2002-11-18
PRIOR APPLICATION NUMBER: US 60/315,315
PRIOR PILING DATE: 2001-08-28
NUMBER OF SEQ ID NOS: 2678
SOFTWARE: Patentin version 3.0
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Sequence 1284, Application US/10230006

Sequence 1284, Application US/10230006

Sequence 1284, Application US/10230006

Sequence 1284, Application US/10230006

GENERAL INFORMATION:

APPLICANT: Robozyme Pharmaceuticals, Inc.

APPLICANT: McSenaugh, Xathy

APPLICANT: McSenaugh, Xathy

TITLE OF INVENTION: METHOD AND REAGENT FOR THE TREATMENT OF ASTHMA AND ALLERGIC CONDI

FILE REFERENCE: 400/056 (MBHB01-1110)
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   PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00669
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00663
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-10-10
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US-10-061-201-716
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US-10-230-006-1249
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
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                                                           2.8%; Score 12.8; DB 1; Length 17; 87.5%; Pred. No. 2.7e+02; tive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
TITLE OF INVENTION: HUMAN POSH-LIKE PROTEIN 1
FILE REFERENCE: PB0178
CURRENT APPLICATION NUMBER: US/10/061,201
CURRENT FILING DATE: 2002-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR PLICATION NUMBER: PCT/US01/00664
PRIOR PLICATION NUMBER: PCT/US01/00664
PRIOR PLING DATE: 2001-01-30
PRIOR PLING DATE: 2001-10-10
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US-10-061-201-716
Sequence 716, Application US/10061201
Publication No. US20030166229A1
GENERAL INFORMATION:
TITLE OF INVENTION: HUMAN POSH-LIKE PROTEIN 1
FILE REPERENCE: PB0178
CURRENT APPLICATION NUMBER: US/10/061,201
CURRENT FILING DATE: 2002-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
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SEQ ID NO 715
LENGTH: 17
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Publication No. US20030166229A1
GENERAL INFORMATION:
                                                                                                                                                                                     226 CTGCCCAGCCCCGAA 241
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                                                        Query Match
Best Local Similarity 87.57
Matches 14; Conservative
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ORGANISM: Homo sapiens
US-10-238-700-3508
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Sequence 751. Application US/10307005
; Publication No. US20030236208A1
; GENERAL INFORMATION:
APPLICANT: University of Delaware
; APPLICANT: University of Delaware
; APPLICANT: Fric B. Kmiec
; APPLICANT: Howard B. Gamper
; APPLICANT: Michael C. Rice
; APPLICANT: Jungub Kim
; TITLE OF INVENTION: Using Modified Single Stranded Oligonuclectides
; TITLE OF INVENTION: Using Modified Single Stranded Oligonuclectides
; TITLE OF INVENTION: Using Modified Single Stranded Oligonuclectides
; TITLE OF INVENTION: Using Modified Single Stranded Oligonuclectides
; TITLE OF INVENTION: Using Modified Single Stranded Oligonuclectides
; TITLE OF INVENTION: Using Modified Single Stranded Oligonuclectides
; FILE REFERENCE: Napro/009 PCT
; CURRENT FILING DATE: 2001-06-01
; PRIOR FILING DATE: 2001-06-01
; PRIOR FILING DATE: 2000-06-01
; PRIOR FILING DATE: 2000-06-01
; PRIOR FILING DATE: 2000-06-01
; PRIOR FILING DATE: 2001-03-07
; RUBBER OF SEQ ID NOS: 271-27
; NUMBER OF SEQ ID NOS: 271-27
; SOFFWANDER.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 17;
          TITLE OF INVENTION: KIT AND METHOD FOR DETERMINING HLA TYPES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
2.8%; Score 12.8; DB 1; Length 1
Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
                                FILE REFERENCE: 1314OP1174
CURRENT APPLICATION NUMBER: US/10/297,068
CURRENT FILING DATE: 2002-11-27
FRIOR APPLICATION NUMBER: JP 2000-164798
PRIOR FILING DATE: 2000-06-01
NUMBER: OF SEQ ID NOS: 1298
SOFTWARE: PatentIn Ver. 2.1
LENGTH: 17
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APPLICANT: University of Delaware APPLICANT: Eric B. Kmiec APPLICANT: Howard B. Gamper
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   410 CTGAGCTGTGGGACGT 425
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                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
ORGANISM: Artificial Sequence
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Matches 14; Conserv
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ORGANISM: Zea mays
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US-10-307-005-752/c
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Sequence 1829, Application US/10430882

Publication No. US20030203870A1

GENERAL INFORMATION:

APPLICANT: Lawrence Blatt

APPLICANT: James McSwiggen

APPLICANT: Dames McSwiggen

APPLICANT: Bharat ChowTira

APPLICANT: Bharat ChowTira

APPLICANT: Bharat ChowTira

APPLICANT: Peter Habberli

TITLE OF INVENTION: Method and Reagent for the Inhibition of NOGO and NOGO Receptor G

FILE REFERENCE: BHBHBO0-878 BH H (4007/112)

CURRENT APPLICATION NUMBER: US/10/430,882

CURRENT FILING DATE: 2001-04-05

PRIOR FILING DATE: 2001-04-05

PRIOR FILING DATE: 2001-02-09

PRIOR FILING DATE: 2001-02-09

PRIOR FILING DATE: 2000-02-11

PRIOR FILING DATE: 2000-02-11

PRIOR FILING DATE: 2000-02-11

PRIOR FILING DATE: 2000-02-11

PRIOR FILING DATE: 2002-04-03

NUMBER OF SEQ ID NOS: 2617

SOFTWARE: Patentin version 3.0

SEQUENCE: DATE: 1000-02-11

SEQUENCE: DATE: 1000-02-11

SEQUENCE: DATE: 2000-03-11

SEQUENCE: DATE: 2000-03-1
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                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
2.8%; Score 12.8; DB 1; Length 17;
Best Local Similarity 68.8%; Pred. No. 2.7e+02;
Matches 11; Conservative 3; Mismatches 2; Indels
CURRENT APPLICATION NUMBER: US/10/230,006
CURRENT FILING DATE: 2002-11-18
PRIOR APPLICATION NUMBER: US 60/315,315
PRIOR FILING DATE: 2001-08-28
NUMBER OF SEQ ID NOS: 2678
SOFTWARE: Patentin version 3.0
SEQ ID NO 1284
LENGTH: 17
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Publication No. US20030228585A1
GENERAL INFORMATION:
APPLICANT: INOKO, Hidetoshi
APPLICANT: KAGIYA, Taeko
APPLICANT: ICHIHARA, Tatsuo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     KAGIYA, Taeko
ICHIHARA, Tatsuo
Matsumura, Yoshiyuki
MORIYA, Shogo
NISHIDA, Michio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        182 GCTGCTGCCCGTTCG 197
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                                                                                                                                                                                                                                                                                    TYPE: RNA
CORGANISM: Homo sapiens
US-10-230-006-1284
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; ORGANISM: Homo sapiens
US-10-430-882-899
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APPLICANT:
APPLICANT:
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US-10-138 674-8374/c

1 Sequence 8374, Application US/10138674

2 Sequence 8374, Application US/10138674

3 Publication No. US20040077565A1

5 CENREAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: Bacobed, Dam

APPLICANT: Bacobed, Jaim

APPLICANT: Stinchcomb, Dan

APPLICANT: Stinchcomb, Dan

APPLICANT: Scobed, Jaime

TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor

TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor

FILE REFERENCE: MBHEBO-876-N (400/049)

CURRENT APPLICATION NUMBER: US/10/138,674

CURRENT PLING DATE: 2002-05-03

NUMBER OF SEQ ID NOS: 20822

SOFTWARE: Patentin Version 3.0

TEMPLY OF APPLICATION OF APPL
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APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Bavco, Pam
APPLICANT: Bavco, Pam
APPLICANT: StinchComb, Jam
APPLICANT: StinchComb, Jam
APPLICANT: Scobedo, Jaime
APPLICANT: Escobedo, Jaime
APPLICANT: Escobedo, Jaime
ITILE OF INVENTION: Merbind and Reagent for the Treatment of Diseases or Conditions Re
FILE REFERENCE: MBHB00-876-N (400/049)
FILE REFERENCE: MBHB00-876-N (400/049)
CURRENT FILING DATE: 2003-04-11
NUMBER OF SEQ ID NOS: 20822
SOFTWARES PATENTON INVENTION OF SEQ ID NOS: 20822
LENGTH: 17
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2.8%; Score 12.8; DB 1;
Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2;
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; Publication No. US20040102389A1
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            363 GCCGCAGGAAGAGAA 378
                                          363 GCCGCAGGAAGAGGAA 378
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: RNA
CORGANISM: Homo sapiens
US-10-138-674-8374
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US-10-712-672-757/c
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APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: Morrises, Dave
ITILE NETRINGE: 400/075 (MAHBOR) 845-1)
CURRENT APPLICATION NUMBER: US/10/342,902
FILE REFERENCE: 400/075 (MAHBOR) 89/077,478
PRIOR APPLICATION NUMBER: US 09/677,478
PRIOR APPLICATION NUMBER: US 09/631,025
PRIOR PELING DATE: 2000-08-09
PRIOR PELING DATE: 2000-08-09
PRIOR PELING DATE: 2000-08-09
PRIOR PELING DATE: 2000-10-24
PRIOR APPLICATION NUMBER: US 09/696,347
PRIOR PELING DATE: 1994-02-07
PRIOR PELING DATE: 1994-02-07
PRIOR PELING DATE: 1995-05-14
PRIOR FILING DATE: 1995-11-08
PRIOR PELING DATE: 1995-11-
APPLICANT: Michael C. Rice
APPLICANT: Jungsup Kim
TITLE OF INVENTION: Targeted Chromosomal Genomic Alterations in Plants
TITLE OF INVENTION: Targeted Chromosomal Genomic Alterations in Plants
TITLE OF INVENTION: Using Modified Single Stranded Oligonucleotides
FILE REFERENCE: Napro/009 PCT 100.05
CURRENT APPLICATION NUMBER: US/005
CURRENT FILING DATE: 2002-11-62
PRIOR FILING DATE: 2001-06-01
PRIOR FILING DATE: 2000-06-01
PRIOR FILING DATE: 2000-06-01
PRIOR FILING DATE: 2000-10-30
PRIOR PLILING DATE: 2000-10-30
PRIOR PLILING DATE: 2000-10-30
PRIOR PLILING DATE: 2001-03-27
NUMBER OF SEQ ID NOS: 2717
SOFTWARE: Friedman macro Napro4
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2.8%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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Publication No. US20040054156A1
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Draper, Kenneth
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ORGANISM: Hepatitis B virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; ORGANISM: Zea mays
US-10-307-005-752
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APPLICANT: Partice, Lee
APPLICANT: Patrice, Lee
APPLICANT: Renneth, Draper
APPLICANT: Eliabeth, Roberta
TITLE OF INVENTION: OLIGOMOCLECTIDE MEDIATED INHIBITION OF HEPATITIS B VIRUS AND HEPJ
TITLE OF INVENTION: VIRUS REPLICATION
FILLS OF INVENTION: VIRUS REPLICATION
CURRENT APPLICATION NUMBER: US /00/2-03-26
FRIOR APPLICATION NUMBER: US 60/296,876
FRIOR APPLICATION NUMBER: US 60/335,059
FRIOR FILLING DATE: 2001-10-05
FRIOR APPLICATION NUMBER: US 60/335,059
FRIOR PRILING DATE: 2001-12-05
FRIOR PRILING DATE: 2001-12-05
FRIOR PAPLICATION NUMBER: US 60/35,124
FRIOR PLILNG DATE: 2001-03-26
FRIOR FILLING DATE: 2001-03-26
FRIOR FILLING DATE: 2001-03-26
FRIOR FILLING DATE: 2001-03-26
FRIOR FILLING DATE: 2000-03-10
FRIOR FILLING DATE: 2000-03-11
FRIOR FILLING DATE: 2000-03-11
FRIOR FILLING DATE: 2000-03-11
FRIOR FILLING DATE: 2000-03-12
FRIOR FILLING DATE: 2000-03-13
FRIOR FILLING DATE: 2000-03-14
FRIOR FILLING DATE: 2000-03-15
FRIOR FILLING DATE: 2000-03-16
FRIOR FILLING 
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APPLICANT: Lawrence, Blatt
APPLICANT: Dennis, Macejak
APPLICANT: Dennis, Macejak
APPLICANT: Dennis, Macejak
APPLICANT: Dennis, Macejak
APPLICANT: Dennis, Marrissey
APPLICANT: Pamela, Pavco
APPLICANT: Pamela, Pavco
APPLICANT: Partice, Lee
APPLICANT: Renneth, Draper
APPLICANT: Renneth, Roberts
TITLE OF INVENTION: UIRUS REPLICATION
FILE REFERENCE: 400/042US (MBHB02-249-E)
CURRENT APPLICATION NUMBER: US/10/669,841
CURRENT FILING DATE: 2003-09-23
PRIOR PELICATION NUMBER: DCT/US02/09187
PRIOR FILING DATE: 2002-03-26
PRIOR PELICATION NUMBER: US 60/296,876
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ilarity 87.5%; Pred. No. 2.7e+02;
Conservative 0; Mismatches 2; Indele
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              117 AGCGGGGGGAAAAGCC 132
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                                                                                                                                       Pamela, Pavco
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Best Local Similarity
Matches 14; Conserv
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Sequence 1149, Application US/10712672

Publication No. US20040102413A1

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: Chowitza, Bharat

APPLICANT: Chowitza, Bharat

APPLICANT: Stinchcomb, Dan

TITLE OF INVENTION: Method and Reagent for the Inhibition of Telomerase Enzyme

FILE REFERENCE: MBHB00-882-C (400/019)

CURRENT APPLICATION NUMBER: US/10/712,672

CURRENT FILING DATE: 2003-11-13

PRIOR PPLICATION NUMBER: 60/197,769

PRIOR PPLING DATE: 1999-08-31

NUMBER OF SEQ ID NOS: 5586

SOFTWARE: PatentIn version 3.0

SEQ ID NOS: 5586
                        APPLICANT: Chowita, Bharat
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
TITLE OF INVENTION: Method and Reagent for the Inhibition of Telomerase Enzyme
FILE REFERENCE: MBHBO0-882-C (400/019)
CURRENT APPLICATION NUMBER: US/09/12,672
CURRENT APPLICATION NUMBER: US/09/653,225
PRIOR PILING DATE: 2000-03-1
PRIOR PLLING DATE: 2000-04-14
PRIOR PLLING DATE: 2000-04-14
PRIOR PLLING DATE: 2000-04-14
PRIOR PLLING DATE: 2000-04-14
PRIOR PLLING DATE: 1999-08-31
NUMBER OF SEQ ID NOS: 5586
SSOTUM RE: Patentin version 3.0
SEQ ID NO 757
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2.8%; Score 12.8; DB 1;
Best Local Similarity 87.5%; Pred, No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 12.8; DB 1;
Pred. No. 2.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-10-669-841-1053/c ; Sequence 1053, Application US/10669841 ; Publication No. US20040127446A1 ; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Sirna Therapeutics, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 263 GGCCGGGGCTTCTCC 278
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 87.5%;
Matches 14; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16 GCCCTGGGGCTTCTCC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16 GCCTTGAAGCCGCGGT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Homo sapiens
US-10-712-672-757
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                     PRIOR FILING DATE: 2001-03-26
PRIOR APPLICATION NUMBER: US 09/740,332
PRIOR FILING DATE: 2000-132
PRIOR FILING DATE: 2000-07-07
PRIOR FILING DATE: 2000-07-07
PRIOR FILING DATE: 2000-07-07
PRIOR FILING DATE: 2000-02-15
PRIOR FILING DATE: 2000-02-15
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 16207
SOFTWARE: PatentIn version 3.0
SEQ ID NO 6332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 16207
SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ·;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; OTHER INFORMATION: oligonucleotide substrate US-10-669-841-6232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 6348, Application US/10669841
Publication No. US20040127446A1
GENERAL INFORMATION:
APPLICANT: Lawrence, Blatt
APPLICANT: Dennis, Macejak
APPLICANT: Dennis, McSaigen
APPLICANT: Dennis, McSaigen
APPLICANT: David, Morrissey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              208 GGACCTGCGGCGGGTC 223
                                                                                                                                                                                                                                                                                                                                                                                   TYPE: RNA
ORGANISM: Artificial Sequence
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APPLICANT: Patrice, Lee
APPLICANT: Kenneth, Draper
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
NAME/KEY: misc_feature
LOCATION:
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Lawrence, Blatt
APPLICANT: Dennis, Macejak
APPLICANT: Dennis, Macejak
APPLICANT: David, Morrissey
APPLICANT: David, Morrissey
APPLICANT: Patrice, Lee
APPLICANT: Patrice, Lee
APPLICANT: Renneth, Draper
APPLICANT: Moberts
APPLICANT: Moberts
TITLE OF INVENTION: UNING REPLICATION
TITLE OF INVENTION: UNING REPLICATION UNING REPLICATION
TITLE OF INVENTION UNING REPLICATION
THOR FILING DATE: 2001-10-24
PRIOR FILING DATE: 2001-12-05
PRIOR FILING DATE: 2002-03-01
PRIOR FILING DATE: 2002-03-01
PRIOR FILING DATE: 2002-03-01
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PRIOR FILING DATE: 2001-06-08

PRIOR APPLICATION NUMBER: US 60/335,059

PRIOR APPLICATION NUMBER: US 60/337,055

PRIOR APPLICATION NUMBER: US 60/337,055

PRIOR FILING DATE: 2001-12-05

PRIOR PILING DATE: 2002-02-02

PRIOR PELICATION NUMBER: US 60/363,124

PRIOR PELICATION NUMBER: US 60/363,124

PRIOR PELICATION NUMBER: US 09/817,879

PRIOR PELICATION NUMBER: US 09/817,879

PRIOR APPLICATION NUMBER: US 09/611,931

PRIOR APPLICATION NUMBER: US 09/611,931

PRIOR PILING DATE: 2000-02-07

PRIOR PELING DATE: 2000-02-07

PRIOR PELING DATE: 2000-02-07

PRIOR PILING DATE: 2000-02-15

PRIOR PILING DATE: 2000-02-15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                j OTHER INFORMATION: oligonucleotide substrate
US-10-669-841-4018
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 6232, Application US/10669841
Publication No. US20040127446A1
GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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NAME/KEY: misc_feature
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LENGTH: 17
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APPLICANT: McSwiggen, James
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level
TITLE OF INVENTION: Number: 40/10/724,270
CURRENT APPLICATION NUMBER: US 60/18,471
PRIOR APPLICATION NUMBER: US 60/296,249
PRIOR FILING DATE: 2001-09-10
PRIOR FILING DATE: 2001-06-06
PRIOR FILING DATE: 2001-06-06
PRIOR FILING DATE: 2001-06-06
PRIOR FILING DATE: 2001-06-10
PRIOR FILING DATE: 2001-06-29
PRIOR FILING DATE: 2002-09-10
PRIOR FILING DATE: 2002-09-10
PRIOR FILING DATE: 2002-06-06
PRIOR FILING DATE: 2002-06-03
PRIOR FILING DATE: 2003-06-23
PRIOR APPLICATION NUMBER: US 10/693,059
PRIOR FILING DATE: 2003-06-31
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Level
TITLE OF INVENTION: RAS, HERZ and HIV
FILE REFERENCE: 400/046-US (MBHB02-326-A)
CURRENT APPLICATION NUMBER: US/10/724,270
CURRENT PILING DATE: 2003-11-26
PRIOR FILING DATE: 2002-05-29
PRIOR FILING DATE: 2001-09-10
PRIOR FILING DATE: 2001-09-10
PRIOR PILING DATE: 2001-06-06
PRIOR APPLICATION NUMBER: US 60/294,140
PRIOR PILING DATE: 2001-05-29
PRIOR PILING DATE: 2001-05-29
PRIOR PILING DATE: 2001-05-29
PRIOR PILING DATE: 2002-06-10
PRIOR PILING DATE: 2002-06-10
PRIOR PILING DATE: 2002-06-06
PRIOR PILING DATE: 2002-06-06
PRIOR PILING DATE: 2002-06-06
PRIOR FILING DATE: 2002-06-09
PRIOR FILING DATE: 2002-06-06
PRIOR FILING DATE: 2002-06-06
PRIOR FILING DATE: 2002-06-06
PRIOR FILING DATE: 2002-06-06
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 12.8; DB 1;
Pred. No. 2.7e+02;
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Similarity 87.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-724-270-1379
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Best Local S
Matches 14
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APPLICANT: Sandberg, Jennifer
APPLICANT: Sandberg, Jennifer
APPLICANT: Sandberg, Jennifer
APPLICANT: Gordon, Glad
APPLICANT: Gordon, Glad
APPLICANT: Stinchcomb, Dan
TITLE OF INVENTION: NUCLEIC ALD BASED MODULATION OF VASCULAR ENDOTHELIAL GROWTH FACT
TITLE OF INVENTION: NUCLEIC ALD BASED MODULATION OF VASCULAR ENDOTHELIAL GROWTH FACT
TITLE OF INVENTION: NUMBER: US (2007)
TITLE OF INVENTION: NUMBER: US (2007)
FILE REFERENCE: MBH02-325PCT (400/047)
FILE REFERENCE: MBH02-325PCT (400/047)
FRICK APPLICATION NUMBER: US (2005)
FRICK APPLICATION NUMBER: US (2007)

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                                                                                                                                                                                  OTHER INFORMATION: Description of Artificial Sequence: Nucleic Acid
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2.8%; Score 12.8; DB 1; Length 17;
Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2; Indels
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1. Similarity 87.5%; Pred. No. 2.7e+02;
14; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                       NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: oligonucleotide substrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 3415, Application US/10712633
Publication No. US20040220128A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 1379, Application US/10724270 Publication No. US20050080031A1
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                                                                         TYPE: RNA
ORGANISM: Artificial Sequence
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ORGANISM: Homo Sapiens
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Best Local Similarity
Matches 14; Conserva
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US-10-712-633-3415/c
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US-10-724-270-1379/c
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SEQ ID NO 6348
LENGTH: 17
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19-10-72-270-4675, Application US/10724270
19-10-72-270-4675, Application US/10724270
19-10-72-270-4675, Application US/10724270
19-Ublication NO US20050080031A1
19-Dblication No US2005009
19-Dblication No US2005-05-29
19-Dblication No US2005-05-05-05
10-Dblication No US2005-05-05-05
10-Dblication No US2005-05-05
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Sequence 5524, Application US/10724270

Publication No. US20050080031A1

SEQUENCE INFORMATION:

APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: McSwiggen, James

TITLE OF INVENTION: Nucleic Acid Treatment of Diseases or Conditions Related to Levels

TITLE OF INVENTION: RAS, HERZ and HIV

TITLE OF INVENTION: RAS, HERZ and HIV

TITLE OF INVENTION: RAS, HERZ and HIV

CURRENT APPLICATION NUMBER: US/10/724,270

CURRENT FILING DATE: 2003-11-26

PRIOR APPLICATION NUMBER: US 60/296,249

PRIOR APPLICATION NUMBER: US 60/296,249

PRIOR APPLICATION NUMBER: US 60/296,249

PRIOR APPLICATION NUMBER: US 60/294,140

PRIOR PILING DATE: 2001-05-29

PRIOR PILING DATE: 2002-09-10

PRIOR PILING DATE: 2002-09-10

PRIOR PILING DATE: 2002-09-10

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Best Local Similarity 87.5%; Pred. No. 2.7e+02;
Matches 14; Conservative 0; Mismatches 2;
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ORGANISM: Homo sapiens
                               US-10-724-270-4675/c
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SGQUENCE 2187, Application US/10724270

SGQUENCE 2187, Application US/10724270

FULL OF UNDIATION:

SAPELICANT: Sina Therapeutics, Inc.

APPLICANT: MCSwiggen, James

TITLE OF INVENTION: RAS, HERZ and HIV

FILE REFERENCE: 400/046-US (WHBHB02-356-A)

CURRENT PELIGANT: 400/046-US (WHBHB02-356-A)

CURRENT PELIGNO NUMBER: US/10/724,270

FRICR FILING DATE: 2003-11-26

FRIOR PELICATION NUMBER: US 60/296,249

FRIOR PELICATION NUMBER: US 60/296,249

FRIOR FILING DATE: 2001-09-10

FRIOR FILING DATE: 2001-09-10

FRIOR FILING DATE: 2001-09-10

FRIOR FILING DATE: 2001-09-10

FRIOR FILING DATE: 2001-05-29

FRIOR FILING DATE: 2002-05-06

FRIOR FILING DATE: 2002-05-06

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PRIOR APPLICATION NUMBER: US 10/444,853
PRIOR FILING DATE: 2003-05-23
PRIOR FILING DATE: 2003-05-23
PRIOR FILING DATE: 2003-04-16
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 6810
SEQ ID NO 1500
LENGTH: 17
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NUMBER OF SEQ ID NOS: 6810
SOFWARE: PatentIn version 3.0
SEQ ID NO 2187
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US-10-724-270-1500
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PRIOR APPLICATION NUMBER: US 10/157,580

PRIOR FILING DATE: 2002-05-29

PRIOR PELING DATE: 2002-10-23

PRIOR PELING DATE: 2002-10-23

PRIOR APPLICATION NUMBER: US 10/44,853

PRIOR APPLICATION NUMBER: US 10/417,012

PROBLEMATE: 2003-04-16

NUMBER OF SEQ ID NOS: 6810

SEQ ID NO 5524

LENGTH: 17

TYPE: RNA

ORGANISM: Homo sapiens

US-10-724-270-5524

Query Match

2.8%; Score 12.8; DB 1; Length 17;

Best Local Similarity 87.5%; Pred. No. 2.7e+02;

Matches 14; Conservative 0; Mismatches 2; Indels 0; Gaps

Oy 6 GCGGAGGGTGGCTG 21

Db 17 GAGGAGGTGGGCTG 2

Search completed: August 24, 2005, 14:36:14

Job time: 4 secs
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